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(cgn2_6/ptodata/1/ina/6A_COMB.seq:*

(cgn2_6/ptodata/1/ina/6B_COMB.seq:*

(cgn2_6/ptodata/1/ina/ecTuS_COMB.seq:*

(cgn2_6/ptodata/1/ina/pcTuS_COMB.seq:*
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(c) 1993 - 2005 Compugen Ltd.
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US-09-423-233-6
Sequence 6, Application US/09423233
Sequence 6, Application US/09423233
Sequence 6, Application US/09423233
Sequence 6, Application US/09423233
Fatent No. 6372430
GOVERNEAL INFORMATION:
APPLICANT: The GOVERNMENT Of the United States of America as TITLE OF INVENTION: Other Filamentous Fungi
TITLE OF INVENTION: Other Filamentous Fungi
TITLE OF INVENTION: Other Filamentous Fungi
CURRENT APPLICATION UNBERS: US/09/423,233
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 61
SOFTWARE: PatentIn Ver. 2.0
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US-08-986-727-2

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US-09-423-233-2

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US-08-986-727-5

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       1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAAATCTTTGAACGCACA 60
                                                                                                                               61 TTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Beck, James J.
TITLE OF INVENTION: DETECTION OF WHEAT FUNGAL PATHOGENS
TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 5827695artis Corporation Patent Department
STREET: 3054 Cormallis Road
CITY: Research Triangle Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (consensus seguence)
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LOCATION: 1..30
OTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
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ZIP: 20779-2257
COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/905,314A
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NAME: Meigs, J. Timothy
REGISTATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: CGC 1944
TELECOMMUNICATION INFORMATION:
TELEFPANE: (919) 541-8689
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 561 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24, Application US/08905314A Patent No. 5827695
GENERAL INFORMATION:
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INDIVIDUAL ISOLATE: 64452 and
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31..181
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CLASSIFICATION: 435
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ORIGINAL SOURCE:
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NAME/KEY:
LOCATION:
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                                                             GENERAL INCORDATION:
APPLICANT: Lee, May
TITLE OF INVENTION: Production of High Titers of Gibberellins GA4 and GA7
FILE REPERENCE: L02-01NP
CURRENT APPLICATION NUMBER: US/09/645,073
CURRENT FILING DATE: 2000-08-25
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 2293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2204 CGCGGCGCGCCAAGCCGTTAAACCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 2263
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Sequence 7. Application US/09423233

GENERAL INFORMATION:

APPLICANT: The Government of the United States of America as TILLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and TILLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and TILLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 20363-0341WP

CURRENT PAPLICATION NUMBER: US/09/423,233

CURRENT PILING DATE: 2000-06-27

NUMBER OF SEQ ID NOS: 61

SEQ ID NO 7

LENGTH: 310

TYPE: APPLICATION OF THE OF THE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 244.4; DB 3; Length Pred. No. 5e-71; 0; Mismatches 31; Indels
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Sequence 1, Application US/09645073
Patent No. 6287800
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
, ORGANISM: Gibberella fujikuroi
US-09-645-073-1
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88.1%;
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Best Local Similarity
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277; Conserva
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Patent No. 5792611
GENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
ITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
ADDRESSEE: Seaby & MacLean
STREET: 880 Wellington Street, Suite 708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
REFERENCE/DOCKET NUMBER: 1898
TELECOMMULICATION INFORMATION:
TELEPHONE: (613) 232-5815
TELEFAX: (613) 232-5831
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 581
STRANDEDNES: double
TOPOLOGY: linear
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ZIP: K1R 6K7
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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NAME: George A. Seaby
REGISTRATION NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1899
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
TELEFAX: (613) 232-5831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          302 AATACCCGCTGAACTTAA 319
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US-08-652-127C-6
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US-08-652-127C-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 75.2%; Score 240; DB 1; Length 561; Best Local Similarity 88.7%; Pred. No. 8e-70; Matches 282; Conservative 2; Mismatches 27; Indels
                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature

COCATION: 505..561

OTHER INFORMATION: /note= "5' end of large subunit

OTHER INFORMATION: rRNA gene"

US-08-905-314A-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 5
US-08-652-127C-6
Squence 6, Application US/08652127C
Squence 10. 5792611
GENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
ITLE OF INVENTION: DETECTION OF PLANT
ITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
ADDRESSEE: George A. Seaby
STREET: 880 Wellington Street, Suite 708
                                                                                                                                                          "5.8S rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: GEOTSP A. SEADY
REGISTRATION NUMBER: 24,034
OTHER INFORMATION: /note= "ITS 1" FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: Canada
ZIP: KIR 6K7
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
LOCATION: 339.504
OTHER INFORMATION: /note= "ITS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        302 AATACCCGCTGAACTTAA 319
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                                                                        NAME/KEY: misc feature
LOCATION: 182..338
OTHER INFORMATION: /note=
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317 TTGCGCCCGCTAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC 376
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                                                                                                                                                                                                                                                                                                                     TTGCGCCCGCCAGTATTCTGGCGGCCATGCCTGTTCGAGCGTCATTACAACCTCAGGCC 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               377 CCCGGGCTTGGTGGTGGGGATCGGCGAGCCTCCGCGCC-----CGCCGTCCCCTAAA 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TACAGTGGCGGTCCCGCCGCGCAGCTTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAGA 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCCGGGCCTGGCGTTGGGGATCGGCGGAAGCCCCTGCGGGCACAACGCCGTCCCCCAAA 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         243 GCGGCGCCGCCACGCCGTAAAACACCCAACTTCTGAATG-TTGACCTCGAATCAGGTAGG 301
                                                                                                                                                                                                                   3 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA
                                                                                                                                                            12;
                                                                                                        65.7%; Score 209.6; DB 1; Length 583; ilarity 85.5%; Pred. No. 1.1e-59; Conservative 0; Mismatches 34; Indels 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
ADDRESSEE: George A. Seaby
ADDRESSEE: Seaby & MacLean
STREET: 880 Wellington Street, Suite 708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NAMER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIPICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: George A. Seaby
REGIENTATION UNUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: Canada
ZIP: KIR 6K7
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-652-127C-5
; Sequence 5, Application US/08652127C
; Patent No. 5792611
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (613) 232-5815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (613) 232-5831
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
US-08-652-127C-5
     STRANDEDNESS: double
                                linear
                                                                                                     Query Match
Best Local Similarity
Matches 272; Conserv
                           ; TOPOLOGY:
US-08-652-127C-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               434 ACAGCGCGCCCCCCTTAAAACCCCCCCACTTCTGAAAGGTTGACCTCGGATCAGGTAGG 493
                                                                                                                                                                                                                                                                                                                                                                          203 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 262
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                                                                                                                                                                                                                                                                                                                        3 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 62
                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                    10;
                                                                                                                                                                                                                65.8%; Score 210; DB 1; Length 531;
85.2%; Pred. No. 7.6e-60;
live 0; Mismatches 37; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8, Application US/08652127C; Sequence 8, Application US/08652127C; Patent No. 57926H; GENEMATION:

APPLICANT: Richard C. Hamelin

TITLE OF INVENTION: PATHOGEN FUNGI

NUMBER OF SEQUENCES: 10

CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
OPPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: GEOCYE A. Seaby
REGISTRENCE/DOCKET NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION: (613) 232-5815
TELEFAX: (613) 232-5815
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 583
TYPE: nuclei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      302 AATACCCGCTGAACTTAA 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 531
                                                                                                                                                                                                  Query Match
Best Local Similarity 85.2
Matches 271; Conservative
                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                ; TOPOLOGY: linear
US-08-652-127C-7
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331 CC---AGCTTGGTGTTGGGAGCTG-------CAGTCCTGCTGCTGCCCCAA 372
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         373 ATACATTGGCGGTCACGTCGRAGCTTCCATAGCGTAGTAATTTACATATCGTTACTGGTA 432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     211 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     242 AGCGGCGCGCCACGCCGTAAAACACCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              433 ATCGTCGCGGCYACGCCGTTAAAC-CCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 491
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
US-005-314A-19
; Sequence 19, Application US/08905314A
; Patent No. 5827695
; GENERAL INFORMATION:
; APPLICANT: Beck, James J.
; TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION NUMBER OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: No. 5827695artis Corporation Patent Department STREET: 3054 Cormallis Road
; CITY: Research Triangle Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 183.8; DB 1; Length 504;
Pred. No. 3.9e-51;
1; Mismatches 38; Indels 19;
                                                                                                                   end of small subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               end of large subunit
         INDIVIDUAL ISOLATE: R-5106, R-5126, and R-5146
                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: 162..318
LOCATION: 162..318
LOCATIOR:
NAME: NEOWATION: /note= "5.85 rRNA gene"
RAME/KEY: misc_feature
LOCATION: 319..472
OTHER INFORMATION: /note= "ITS 2"
                                (consensus seguence)
                                                                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: 13..161
OTHER INFORMATION: /note= "ITS 1"
                                                                     NAME/KEY: misc_feature
LOCATION: 1.12
THER INFORMATION: /note= "3'
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "5'
rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:

NAME/KEY: misc_feature

LOCATION: 473.504

OTHER INFORMATION: /note=

COTHER INFORMATION: rNA give-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 57.6%;
Best Local Similarity 81.5%;
Matches 255; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          302 AATACCCGCTGAA 314
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                             INDIVIDUAL ISOLATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      122
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                                                                                                                   TIGGGCCCCCCAGIATICIGGCGGCCATGCCTGTICGAGCGTCATTACAACCCTCAGGCC 122
                                                                                                                                                                                                                                                                      123 CC----CGGGCCTGGCGTTGGGGATCGGCGGAAGCCCCTGCGGGCACACGCCGTCCCC 178
                                                                                                                                                                                                                                                                                                                                                                                         318 TTGCGCCCCCCAGTATTCTGCCGGCCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCA 377
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 239 GAGAGCGGCGCCGCCACGCCGTAAAACACCCAACTTCTGAAT-GTTGACCTCGAATCAGG 297
                                                                                              62
                                                                                         3 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA
                                                    Gaps
                                                  10;
Query Match
60.1%; Score 191.6; DB 1; Length 594;
Best Local Similarity 83.2%; Pred. No. 1.1e-53;
Matches 268; Conservative 0; Mismatches 44; Indels 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Beck, James J
TITLE OF INVENTION: Detection of Fungal Pathogens Using the
TITLE OF INVENTION: Polymerase Chain Reaction
NUMBER OF SEQUENCES: 96
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REPERENCE/DOCKET NUMBER: CGC 1739/PCT/CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
TELEFAX: 919-541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ...urkESSEE: No. 5814453artis Corporation STREET: 520 White Plains Road CITY: Tarrytown STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/722,187
FILING DATE: 15-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        298 TAGGAATACCCGCTGAACTTAA 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    553 TÁGGÁCTÁCCCGCTGAACTTAA 574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10591
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 82, Application US/08887480 Patent No. 5814453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Fusarium culmorum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 504 Base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-887-480-82
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Gaps

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/note= "DNA sequence for the internal transcribed spacer region of Fusarium culmorum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 ATTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGC 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ----cadrecrecrecacrecea 372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62 ATTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTACAACCCTCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122 CCCCGGGCCTGGCGTTGGGGATCGGCGGAAGCCCCTGCGGGCACAACGCCGTCCCCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 504;
                                                                                                                                                                                                                                                      APPLICANT: Ligon, James M
APPLICANT: Ligon, James M
TITLE OF INVENTION: Detection of Fungal Pathogens Using the
TITLE OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
ADDRESSE: Ciba-Geigy Corporation
STREET: 7 Skyline Drive
CITY: Hawthorne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION OPPLAR: US 08/233,608
FILING DATE: 04-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 57.6%; Score 183.8; DB 2; Best Local Similarity 81.5%; Pred. No. 3.9e-51; Matches 255; Conservative 1; Mismatches 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                              Sequence 82, Application US/08722187
Patent No. 5955274
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (fculm.con)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Walsh, Andrea C.
REGISTRATION NUMBER: 34,988
REPERENCE/DOCKET NUMBER: GG TELECOMMUNICATION INPORMATION:
TELEFAX: 919-541-8666
TELEFAX: 919-541-8669
INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
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LOCATION: 1..504

OTHER INFORMATION: /note:

OTHER INFORMATION: inter:

OTHER INFORMATION: (fculi
     302 AATACCCGCTGAA 314
                                                        492 AATACCCGCTGAA 504
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                   RESULT 11
US-08-722-187-82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   271 ATTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGC 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62 ATTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTACAACCCTCAGGC 121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               331 CC---AGCTTGGTGTTGGGAGCTG-------CAGTCCTGCTGCACTCCCAA 372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATACAGTGGCGGTCCCGCCGCAGCTTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAG 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGCGGCGCCGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATGGTGGGGGCYACGCCGTTAAAC-CCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  373 ATACATTGGCGGTCACGTCGRAGCTTCCATAGCGTAGTAATTTACATATCGTTACTGGTA 432
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc feature
LOCATION: 1..12
OTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: 473..50
OTHER INFORMATION: /note= "5' end of large subunit
OTHER INFORMATION: FRNA gene"
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/905,314A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Fusarium culmorum
INDIVIDUAL ISOLATE: R-5106, R-5126, and R-5146
INDIVIDUAL ISOLATE: (consensus sequence)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "5.8S rRNA gene'
                                                                                                                                                                      ATTORNEY AGENT INFORMATION:
NAME: Meigs, J. Timothy
REGISTAATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: CGC 1944
TELECOMMUNICATION INFORMATION:
TELEPAX: (919) 541-8687
TELEPAX: (919) 541-8687
TELEPAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 504 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: 13..161
OTHER INFORMATION: /note= "ITS
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LOCATION: 319.7472
OTHER INFORMATION: /note= "ITS
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LOCATION: 162.7318
OTHER INFORMATION: /note=
                                                                                                                              FILING DATE:
CLASSIFICATION: 435
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182 ATACAGTGGCGGTCCCGCCGCAGCTTCCATTGCGTAGCTAACACCCTCGCAACTGGAG 241
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                                                                                                                                                                                                                                                     242 AGCGGCGCGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
                                                                                                                                                                                                                                                                                CCCCGGGCCTGGCGTTGGGGATCGGCGGAAGCCCCTGCGGGCACAACGCCGTCCCCCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 84, Application US/08887480
Patent No. 5814453
PAPLICANT: Beck, James J
TITLE OF INVENTION:
TITLE OF INVENTION:
POlymerase Chain Reaction
NUMBER OF SEQUENCES: 96
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LOCATION: 1..30
OTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10591
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 96
CORRESPONDENCE ADDRESS:
ADDRESSE: No. 5814453art18 Corporation
STREET: 520 White Plains Road
CITY: Tarrytown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US 08/722,187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 15-0CT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy
REGIGSTRATION NUMBER: 38, 241
REFERENCE/DOCKET NUMBER: GGC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
ORGANISM: Fusarium monilife
                                                                                                                                                                                                                                                                                                                                                   AATACCCGCTGAA 314
                                                                                                                                                                                                                                                                                                                                                                                            492 AATACCCGCTGAA 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 545 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-887-480-84
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LOCATION: 1..504
OTHER INFORMATION: /note= "DNA sequence for the
OTHER INFORMATION: internal transcribed spacer region of Fusarium culmorum
OTHER INFORMATION: (fculm.con)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          270
182 ATACAGTGGCGGTCCCGCCGCAGCTTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAG 241
                                      373 ATACATTGGCGGTCACGTCGRAGCTTCCATAGCGTAGTAATTTACATATCGTTACTGGTA 432
                                                                                      242 AGGGGGGGGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
                                                                                                                      433 ATCGTCGCGGCYACGCCGTTAAAC-CCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62 ATTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTACAACCCTCAGGC 121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 57.6%; Score 183.8; DB 5; Length 504; Best Local Similarity 81.5%; Pred. No. 3.9e-51; Matches 255; Conservative 1; Mismatches 38; Indels 19
                                                                                                                                                                                                                                                                                                                  PCT-US95-04712-82
Sequence 82, Application PC/TUS9504712
GENERAL INFORMATION:
APPLICANT: Ligon, James M
APPLICANT: Ligon, James J
TITLE OF INVENTION: Detection of Fungal Pathogens Using the TITLE OF INVENTION: Polymerase Chain Reaction
NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10522

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRPLICATION:
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/233,608
FILING DATE: 04-ARR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Walsh, Andrea C.
REGISTRATION NUMBER: 34,988
REFERENCE/DOCKET NUMBER: CGC 1739
TELEFONMUNICATION INFORMATION:
TELEFONMUNICATION INFORMATION:
TELEFONS: 919-541-8666
TELEFAX: 919-541-8666
TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
TENTILE SO4 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Ciba-Geigy Corporation
STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                               302 AATACCCGCTGAA 314
                                                                                                                                                                                                                           492 AATACCCGCTGAA 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 504 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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RESULT 15
US-08-887-480-96
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                                                                                                                                                                                                                                                                                                                                                                               56.6%; Score 180.4; DB 1; Length 545; 82.4%; Pred. No. 5.5e-50; tive 0; Mismatches 36; Indels 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: BECK, James J.
TITLE OF INVENTION: DETECTION OF WHEAT FUNGAL PATHOGENS
TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 5872695artis Corporation Patent Department
STREET: 3054 CORNWAllis Road
CITY: Research Triangle Park
                                                                                                                                                                                                                                                 NAME/KEY: misc_feature;
CCATION: 489..545
COTHER INFORMATION: /note= "5' end of large subunit;
COTHER INFORMATION: rRNA gene"
US-08-887-480-84
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APPLICATION NUMBER: US/08/905,314A FILIGN DATE:
CLASSIFICATION: 435
                                                                                                                         "5.8S rRNA gene"
                                                                                                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION: 336..488
OTHER INFORMATION: /note= "ITS 2"
NAME/KEY: misc feature
LOCATION: 31..178
OTHER INFORMATION: /note= "ITS 1"
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ZIP: 20779-2257
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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LOCATION: 179..335
OTHER INFORMATION: /note=
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Best Local Similarity 82.4
Matches 262; Conservative
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449 ATCGTCGCGGCCACGCCGTTAAAC-CCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 507
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: 1..30
OTHER INFORMATION: /note= "3' end of small subunit
FEATURE: INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
LOCATION: 489..545
COTHER INFORMATION: /note= "5' end of large subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "5.8S rRNA gene"
NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: CGC 1944
TELEPHONE: (919) 541-8587
TELEPHONE: (919) 541-8689
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 545 base pairs
TYPE: nucleic acid
STRANDENNESS: single
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LOCATION: 31..178
OTHER INFORMATION: /note= "ITS 1"
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Fusarium moniliforme INDIVIDIAL ISOLATE: 4551
IMMEDIATE SOUNCE:
CLONE: pCRFMON1
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MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
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LOCATION: 336._488
OTHER INFORMATION: /note=
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LOCATION: 179..335
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Sequence 96, Application US/0883480

GENERAL ANTO BACK.

THILE OF INVENTION: Defection of Fungal Pathogens Using the TITLE OF INVENTION: Defection of Fungal Pathogens Using the TITLE OF INVENTION: Defection of Fungal Pathogens Using the TITLE OF INVENTION: Defection of Fungal Pathogens Using the TITLE OF INVENTION: Defection of STREET: 30 Maire Plains Road CITY: Tarrytown STATE: 320 Maire Plains Road CITY: Tarrytown STATE: NW COMPUTES IN PROCEED COMPUTES IN PROCEDUTES IN PROCEED COMPUTES IN PROCEDUTES IN PROCEED COMPUTED IN TAKE IN PROCEDUTES IN PROCEED IN PROCEDUTES IN PROCEDUTES IN PROCEDUTES IN PROCEDUTES IN PROCEED IN PROCEED IN PROCEDUTES IN PROCEDUTES IN PROCEED IN PROCEDUTES IN PROCEED IN PROC
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56.4%; Score 180; DB 1; Length 546; 82.4%; Pred. No. 7.5e-50;

Query Match Best Local Similarity

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290 ATTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGC 349
                                                                                                                                                                                                                                182 ATACAGTGGCGGTCCCGCCGCAGCTTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAG 241
                                                         230 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC 289
                                                                                                                                                    CCCCGGGCCTGGCGTTGGGGATCGGCGGAAGCCCCTGCGGGCACAACGCCGTCCCCAA 181
                                                                                         62 ATTGCGCCCGCCAGTATTCTGGCGGCCATGCCTGTTCGAGCGTCATTACAACCCTCAGGC
21; Gaps
0; Mismatches 35; Indels
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Matches 262; Conservative
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October 23, 2005, 15:20:46 ; Search time 1379.23 Seconds (without alignments) 1907.926 Million cell updates/sec
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(cgn2_6/ptodata/2/pubpna/USO7_NEW_PUB.seq:*
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(cgn2_6/ptodata/2/pubpna/USO7_NEW_PUB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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319
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	Description	Seguence 6,	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
SUMMARIES	QI	US-10-046-955-6	US-10-829-661-5	US-10-672-300-5	US-09-961-755A-7	US-10-773-904-7	US-10-773-905-7	US-10-046-955-7
	BB :	15	21	21	10	20	20	15
	Query Match Length DB	319	552	552	534	534	534	310
æ	Query Match	100.0	81.2	81.2	75.4	75.4	75.4	75.3
	Score	319	259	259	240.4	240.4	240.4	240.2
	Result No.	1	8	e	4	S	9	7

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-09-961-755A	-T96-60-80 0	US-10-773	0 US-10-773-	0 US-10-773-	0 US-10-773-	0 US-09-961-	0 US-10	0 US-10	US-10	US-10	US-10-623-432-	US-11	US-10-046-955-	US-10-757-093-	US-10-046-955-	US-10-046-955-	US-10-757-09	US-10-121-7	ᅻ	US-11-131-659		1 US-1	-994-60-SD	3-1	0S-09		US-10-046-955-	15 US-10-046-955-25		7 US-10-356-320-	5 US-10-046-95	US-09-961-66	0S-09	US-09-961-66	US-09-961-66	-10-829-6	21 US-10-672-300-6
522	222	522	225	522	522	521	521	521	780	652	652	652	365	577	365	364	528	650	650	650	296	296	641	346	618	343	344	346	309	466	336	377	377	377	377	495	495
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ALIGNMENTS

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301 GAATACCCGCTGAACTTAA 319
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US-10-672-300-5
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Best Local Similarity
Matches 295; Conserv
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US-09-961-755A-7
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                                                                                                                                                                           1 GAAAATGCGATAAGTAATGTGAATTGCAGAATTCAGGGAATCATCGAATCTTTGAACGCA 60
                                                                                                                   GAAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCA
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     Length 319;
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APPLICANT: HAN, XTANG-YANG
APPLICANT: TARRAND, JEFREY J.
APPLICANT: PHAM, AUDREY S.
TITLE OF INVENTION: DIAGNOSIS OF MOULD INFECTION
FILE REFERENCE: UTSCREGUE
CURRENT APPLICATION NUMBER: US/10/829,661
PRIOR APPLICATION NUMBER: 10/672,300
PRIOR PILING DATE: 2003-09-26
PRIOR FILING DATE: 2003-09-26
PRIOR FILING DATE: 2003-09-26
PRIOR FILING DATE: 2003-09-27
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PARENTIN Ver: 2.1
SEQ ID NO 5
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  ; Score 319; DB 15;
; Pred. No. 4.9e-103;
0; Mismatches 0;
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     100.0%;
100.0%;
Query Match
Best Local Similarity 100.
Matches 319; Conservative
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ORGANISM: Fusarium sp
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US-10-829-661-5
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414 TTCAGTGGCGGTCTCGCTGCAGCCTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAAG 473
                                                                                                                                          474 GCGGCGCGCCCACGCCGTAAAACCCCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAG 533
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Publication No. US20030113722A1
GENERAL INFORMATION:
APPLICANT: Back, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Delection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Delection of Fusarium Reaction
FILLS REPRENSIVE: 60055
CURRENT APPLICATION WHORER: US/09/961,755A
CURRENT FILING DATE: 2001-09-24
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Sequence 5, Application US/10672300

Publication No. US200S0048509A1

GENERAL INFORMATION:
APPLICANT: HAN, XIANG-YANG

APPLICANT: TARRAND, JEFFREY J.

TITLE OF INVENTION: DIAGNOSIS OF INVASIVE MOLD INFECTION
FILE REFERENCE: UTXC.766US

CURRENT APPLICATION NUMBER: US/10/672,300

CURRENT FILING DATE: 2003-09-26

PRIOR FILING DATE: 2002-09-27

NUMBER OF SEQ ID NOS: 6

SOFTWARE: Patentin Ver. 2.1
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Similarity
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Matches 276;
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US-10-773-905-7
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US-10-046-955-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the TITLE OF INVENTION: Polymerase Chain Reaction
TITLE OF INVENTION: Polymerase Chain Reaction
CURRENT APPLICATION NUMBER: US/10/773,904
PRIOR APPLICATION NUMBER: US/09/961,755
RIGHT APPLICATION NUMBER: US/09/961,755
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                 2 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC
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                                                                                                                                                                  31; Indels
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                                                                                                                                 Score 240.4; DB 10
Pred. No. 6.8e-75;
0; Mismatches 31
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Pred. No. 6.8e-75;
0; Mismatches 31
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Publication No. US20040259120A1
GENERAL INFORMATION:
APPLICANT: Beck, Jim
                                                                               ORGANISM: Fusarium proliferatum
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NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn version 3.0
                                                                                                                                 Query Match 75.4%;
Best Local Similarity 87.9%;
Matches 276; Conservative
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Best Local Similarity 87.9%;
Matches 276; Conservative
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US-10-773-904-7
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                                SEQ ID NO 7
                                                                   TYPE: DNA
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                                                LENGTH:
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Sequence 7, Application US/10046955;
Publication No. US20030129600A1
GENERAL INFORMATION:
APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: Secretary of the Department of Health and Human Services, Centers for Die
                                                                                                           461 CGCGGGGCCAAGCCGTTAAAACCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 520
348 CCCCGGGTTTGGTGTTGGGGATCGGCGAGCCCTTGCGGC-----AAGCCGGCCCCGAA 400
                                                                                                                                                                                                     242 AGCGGCGCGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
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Publication No. US20040259121A1
GENERAL INFORMATION:
APPLICANT: Beck, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Dolymerase Chain Reaction
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/10/773,905
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTHARE: Patentin version 3.0
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; ORGANISM: Fusarium proliferatum
US-10-773-905-7
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                                                                               TITLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and TITLE OF INVENTION: Other Filamentous Fungi
FILE REPERENCE: 6395-6206, 100 CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US/10/046,955
CURRENT FILING DATE: 2000-06-27
PRIOR APPLICATION NUMBER: US 09/423,233
PRIOR APPLICATION NUMBER: PCT/US98/08926
PRIOR PILING DATE: 1998-06-21
PRIOR PELING DATE: 1998-06-01
PRIOR PELING DATE: 1997-05-02
NUMBER OF SEQ ID NOS: 61
SEQ ID NOS: 61
SEQ ID NOS: 61
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APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Delymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/09/961,755A
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
SEQ ID NO 5
LENGTH: 522
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87.4%; Pred. No. 6.6e-75;
live 0; Mismatches 33
Control and Prevention
Morrison, Christine J.
Reiss, Errol
Aidorevich, Liliana
Choi, Jong Soo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09961755A; Publication No. US20030113722A1; GENERAL INFORMATION:
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ORGANISM: Fusarium subglutinans
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US-09-961-755A-5
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US-09-961-755A-5
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US-10-046-955-7
              APPLICANT:
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APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Delection of Fusarium Species infecting Corn Using the
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/09/961,755A
CURRENT PILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
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                                                                                                                                             288 ATTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGC
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                                                                                                                  2 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC
                                                                                                                                                                                                                               62 ATTGCGCCCGCCAGTATTCTGGCGGCCATGCCTGTTCGAGCGTCATTACAACCCTCAGGC
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DB 10; Length 522;
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                                                          Indels
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US-09-961-755A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 57.4%; Score 183.2; DB 10; Best Local Similarity 83.4%; Pred. No. 1.8e-54; Matches 262; Conservative 0; Mismatches 33;
Score 183.2; DB 1
Pred. No. 1.8e-54;
0; Mismatches 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 8, Application US/09961755A; Publication No. US20030113722A1; GENERAL INFORMATION:
  Query Match 57.4%;
Best Local Similarity 83.4%;
Matches 262; Conservative
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US-09-961-755A-8
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LENGTH: 522
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182 ATACAGTGGCGGTCCCGCCGCAGCTTCCATTGCGTAGTAGCTAACACCTCGCAACTGGAG 241
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                                                                                              Query Match
Best Local Similarity 83.4%;
Matches 262; Conservative
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US-10-773-905-5
      LENGTH: 522
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APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT PAPLICATION NUMBER: US/10/773,904
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn version 3.0
SEQ ID NO 5
LENGTH: 522
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Publication No. US20040259120A1
GENERAL INFORMATION:
APPLICANT: Beck, Jim
APPLICANT: Beck, Jim
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/10/773,904
PRIOR FILING DATE: 2004-02-06
PRIOR FILING DATE: 2001-09-24
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83.4%; Pred. No. 1.8e-54;
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                                                                                                                              Sequence 5, Application US/10773904 Publication No. US20040259120A1 GENERAL INFORMATION:
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SOFTWARE: Patentin version 3.0
SEQ ID NO 8
302 AATACCCGCTGAAC 315
                  AATACCCGCTGAAC 315
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Matches 262; Conservative
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APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
FILE REFERENCE: 60055
CURRENT APPLICANTION NUMBER: US/10/773,905
CURRENT FILING DATE: 2004-02-06
PRIOR FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFFWARE: Patentin version 3.0
SEQ ID NO 5
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                                                                                                                Score 183.2; DB 20; Length 522;
Pred. No. 1.8e-54;
0; Mismatches 33; Indels 19;
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TYPB: DNA
ORGANISM: Fusarium verticillioides (syn. F. moniliforme)
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US-09-961-755A-6
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US-10-773-904-6
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Publication No. US20030113722A1
GENERAL INFORMATION:
APPLICANT: Beck, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
AGCGGCGCGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
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Publication No. US20040259121A1
GENERAL INFORMATION:
APPLICANT: Beck, Jim
APPLICANT: Bernett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFRENCE: 60055
CURRENT PAPLICATION NUMBER: US/09/961,773,905
CURRENT PILING DATE: 2004-02-06
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR PILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
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ORGANISM: Fusarium verticillioides (syn. F. moniliforme)
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Pred. No. 1.8e-54;
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Best Local Similarity 83.4%;
Matches 262; Conservative (
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APPLICANT: Barnett, Jason
APPLICANT: Barnett, Jason
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/10/773,904
CURRENT FILING DATE: 2004-02-06
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTMARE: Patentin version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   122 ccccedeccredecerregearcesceaagccccrecesecacacacccccaa 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    242 AGCGGCGCGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGG 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             449 ATCGTCGCGGCCACGCCGTTAAAC-CCCAACTTCTGAATGTTGACCTCGGATCAGGTAGG 507
                                                                                                                                                                                                                                                                                                                                                                                                   228 AAAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCAC 287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 348 CC---AGCTTGGTGTTGGGAGCTG------CAGTCCTGCTGCTGCACTCCCAA 389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     182 ATACAGTGGCGGCCCCCCCCCCCATCCATTGCGTAGTAGCTAACACCTCGCAACTGGAG 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     390 ATACATTGGCGGTCACGTCG-AGCTTCCATAGCGTAGTAATTTACACATCGTTACTGGTA 448
                                                                                                                                                                                                                                                                                                                                                                                                                                                       62 ATTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTACAACCCTCAGGC
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.4.8e-52;
36;
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0; Mismatches 36;
CURRENT APPLICATION NUMBER: US/09/961,755A CURRENT FILING DATE: 2001-09-24 NUMBER OF SEQ ID NOS: 24 SEQ TWARE: Patentin version 3.0 SEQ ID NO 6 LENGTH: 521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 6, Application US/10773904; Publication No. US20040259120A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 82.2%;
Matches 258; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          302 AATACCCGCTGAAC 315
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                                                                                                                                                            TYPE: DNA ORGANISM: Gibberella zeae
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US-10-773-904-6
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        122 CCCCGGGCCTGGCGTTGGGGATCGCGGGAAGCCCCTGCGGGCCAAAGGCCCCCCAA 181

        Db
        348 CC--AGCTTGGGAGCTGCCTGCTGCTGCTGCTGCTCCCCCAA 389

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        182 ATACAGTGGCGCCCCCGCCGCGCGACTTGCGTACTAACTCTGCTAACTGCAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGCGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAACTGAAC
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/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/RECOMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
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               Appli
                                                                                                                                                                                                                                                                                                             Sequence 7, Application US/09423233
Patent No. 6372430
GENERAL INFORMATION:
APPLICANT: The Government of the United States of America as APPLICANT: The Government of the United States of America as TITLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and TITLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 03063-0314WP
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 61
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC
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100.0%; Pred. No. 5.7e-94;
ive 0; Mismatches 0;
US-08-742-023-7

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US-08-233-608-47

US-08-237-990B-7

US-08-123-187-47

US-08-722-187-47

US-09-517-790-85

US-09-517-790-85

US-08-905-314A-23

US-08-905-314A-23

US-08-905-314A-23

US-08-968-505-6

US-08-968-505-6
                                                                                                                                                                                                                                                       ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Fusarium moniliforme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 310; Conservative
  CTGAACTTAA 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGAACTTAA 310
  LENGTH: 310
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1224.4
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1234.4
1233.6
1233.6
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136
131.4
126.6
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RESULT 2 US-09-645-073-1

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                                                                                                                     Production of High Titers of Gibberellins GA4 and GA7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Beck, James J.
TITLE OF INVENTION: DETECTION OF WHEAT FUNGAL PATHOGENS
TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION
TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 5827695artis Corporation Patent Department
STREET: No. 5827695artis Corporation Patent Department
STREET: NC
COUNTRY: NCS
COUNTRY: USA
ZIP: 20779-2257
COMPUTER: READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 306.8; DB 3 Pred. No. 1.6e-92;
Sequence 1, Application US/09645073
Patent No. 6287800
GENERAL INFORMATION:
APPLICANT: Lee, May
APPLICANT: Lee, May
TITLE OF INVENTION: Production of High Tite
CURRENT APPLICATION NUMBER: US/09/645,073
CURRENT FILING DATE: 2000-08-25
PRIOR PILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 1
SOFTWARE: Patentin version 3.0
SEQ ID NO 1
LENGTH: 2293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-905-314A-24; Sequence 24, Application US/08905314A; Patent No. 5827695; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Gibberella fujikuroi
                                                                                                                                                                                                                                                                                                                                                                                                                                      99.0%;
99.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 99.4
Matches 308; Conservative
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                                                                                                                                                                                                                                                                                                                           STRAIN: Fusarium avenaceum
INDIVIDUAL ISOLATE: 64452 and R-4045 (consensus sequence)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: 505..561
OUTHER INFORMATION: /note= "5' end of large subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                     end of small subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match 80.6%; Score 249.8; DB 1; Best Local Similarity 91.6%; Pred. No. 1.1e-73; Matches 285; Conservative 1; Mismatches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "5.8S rRNA gene"
             NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: GGC 1944
TELEPHONE: (919) 541-8587
TELEPHONE: (919) 541-8689
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 561 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: 31..I81
OTHER INFORMATION: /note= "ITS 1"
                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: 1..30
OTHER INFORMATION: /note= "3'
OTHER INFORMATION: rRNA gene"
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                                                                                                                                                                                                                                        AUGUCOTY: linear MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: 182.338
OTHER INFORMATION: /note=
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LOCATION: 339. ..504
OTHER INFORMATION: /note=
ATTORNEY/AGENT INFORMATION:
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240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   241 GGCCAAGCCGTTAAACCCCCAACTTCTGAATG-TTGACCTCGGATCAGGTAGGAATACCC 299
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                                                                                                                                                                                                                                                                                                                                                     Query Match 74.5%; Score 231; DB 1; Length 583; Best Local Similarity 90.4%; Pred. No. 2.3e-67; Matches 281; Conservative 0; Mismatches 25; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 7, Application US/08652127C
| Patent No. 5792611
| GENERAL INFORMATION: APPLICANT: Richard C. Hamelin TITLE OF INVENTION: DETECTION OF PLANT; TITLE OF INVENTION: DETECTION OF PLANT; TITLE OF INVENTION: DETECTION OF PLANT; NUMBER OF SEQUENCES: 10
| CORRESPONDENCE ADDRESS: ADDRESSEE: George A. Seaby; ADDRESSEE: Seaby & MacLean; STREET: 880 Wellington Street, Suite 708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
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ZIP: KIR 6K7
COMPUTER READABLE FORM:
COMPUTER: 3.5 inch diskette
COMFUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
        REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
TELEPHONE: (613) 232-5815
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
TELEFAX: (613) 232-5815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: George A. Seaby
REGISTRATION NUMBER: 24,0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 300 GCTGAACTTAA 310
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                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ottawa
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US-08-652-127C-8
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US-08-652-127C-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     183 TACAGIGGGGGCCGCGCGCGCTICCATIGGGTAGTAGCTAACACCTCGCAACIGGAGA 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCGGCGCGCCAAGCCGTTAAACCCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGA 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      243 GCGGCGCGCCACGCCGTAAAACACCCCAACTTCTGAATGTTGACCTCGAATCAGGTAGGA 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 6372430
GENERAL INFORMATION:
APPLICANT: The Government of the United States of America as
APPLICANT: The Government of the United States of America as
TITLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and
TITLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 03063-0341WP
CURRENT APPLICATION NUMBER: US/09/423,233
CURRENT PILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTACAACCCTCAGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATCTTTGAACGCACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 319;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           33; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 77.5%; Score 240.2; DB 3; Best Local Similarity 87.4%; Pred. No. 1.5e-70; Matches 277; Conservative 0; Mismatches 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
COUNTRY: Canada
ZIP: KIR 6K7
COMPUTER: BM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: BM SCII
CURRENT ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 8, Application US/08652127C Patent No. 5792611
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              294 ATACCCGCTGAACTTAA 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    303 ATACCCCCTCAACTTAA 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: George A. Seaby
REGISTRATION NUMBER: 24,034
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Fusarium solani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                          319
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                                                                                                                                                                                                                                                                                         SEQ ID NO 6
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299 CGCTGAACTTAA 310
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STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TRY: Canada
KIR 6K7
                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
       STRANDEDNESS:
                      TOPOLOGY:
US-08-652-127C-6
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US-08-652-127C-5
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                                                                                                                                                                                                                                                                                            61 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC 120
                                                                                                                                                                                                                                                                                                                            263 TIGCGCCCGCCAGIAIICIGGCGGGCAIGCCIGIICGAGCGICAIIICAACCCICAAGCC 322
                                                                                                                                                                                                                                                                                                                                                                  CCCGGGTTTGGTGTTGGGGATCGGCAAG-CCCTTGCGGCAAGCCGGCCCCGAAATCTAGT 179
                                                                                                                                                                                                                                                                                                                                                                                                                                         GGCGGTCTCGCTGCAGCTTCCATTGCGTAGTAGAAAACCCTCGCAACTGGTACGCGGCG 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGCGGTCTCGCTGTAGCTTCCTCTGCGTAGTAGCACACCTCGC---ACTGGAAAACAGCG 439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CGGCCAAGCCGTTAAACCCCCAACTTCTG-AATGTTGACCTCGGATCAGGTAGGAATACC 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATCTTTGAACGCACA
                                                                                                                                                                                     Gaps
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                                                                                                                                              Score 225.2; DB 1; Length 531;
Pred. No. 2e-65;
0; Mismatches 30; Indels 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: George A. Seaby
E: Seaby & MacLean
880 Wellington Street, Suite 708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWATING STATES ASCII
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: George A. Seaby
REGISTRATION NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (613) 232-5815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 6, Application US/08652127C
Patent No. 5792611
                                                                                                                                                72.6%;
88.8%;
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MEDIUM TYPE: 3.5 inch opponing
                                                                                                                                          Query Match
Best Local Similarity 88.8°
Matches 277; Conservative
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 531
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    500 cecreaacriaa 511
                                                                   ESS: double
linear
                                                  ; TYPE: nucleic acid
; STRANDEDNESS: doubl
; TOPOLOGY: linear
US-08-652-127C-7
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STREET: 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-652-127C-6
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TYPE: n
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253 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 312
                                                                                                                                                                                                                                                                                                490 TGGCCACGCCGTTAAACCCCCCACTTCTGAAAGGTTGACCTCGGATCAGGTAGGAATACC 549
                                                                                                                                                                                                                                                            TIGCGCCCCCCAGIAITCIGGCGGGCAIGCCIGITCGAGCGICAITICAACCCICAAGCC 120
                                                                                                                                                                                                                                                                                                                                                                                          121 CCCGGGTTTGGTGTTGGGGATCGGCAAG-CCCTTGCGGCAAGCCGGCCCCGAAATCTAGT 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      433 GGCGGTCTCGCTGTAGCTTCCTCTGCGTAGTAGCACACCTCGC---ACTGGAAAACAGCG 489
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                                                                                                                                 1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATCTTTGAACGCACA
                                                                  5,
      Length 581;
                                                                  Indels
Query Match 72.3%; Score 224.2; DB 1; Best Local Similarity 88.5%; Pred. No. 4.4e-65; Matches 276; Conservative 1; Mismatches 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: George A. Seaby
ADDRESSEE: Seaby & MacLean
STREET: 880 Wellington Street, Suite 708
CITY: Ottawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SIGLE.
SOFTWARE. ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGBTT INFORMATION:
NAME: George A. Seaby
REGISTRATION NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/08652127C
Patent No. 5792611
GENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
TITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IEM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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61 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C---AGCTTGGTGTTGGG-----ATCTGTGCAAACACAGTCCCCAAATTGATTG 398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        399 GCGGTCACG-TCGAGCTTCCATAGCGTAGTAATTTACACATCGTTACTGGTAATCGTCGC 457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                241 GGCCAAGCCGTTAAACCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      458 GGCCACGCCGTTAAA-CCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       231 AAATGCGATAAGTAAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA
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US-08-905-314A-22
iS-quence 22, Application US/08905314A

Patent No. 5827695
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
USING THE POLYMERASE CHAIN REACTION
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEB: No. 5827695artis Corporation Patent Department
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: NC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60.5%; Score 187.6; DB 1; Length 546; 84.5%; Pred. No. 8.1e-53; ive 0; Mismatches 34; Indels 14
        T-427, T-534, and T-756 (consensus
                                                              CLONE: pCRFpoaeT427(1-2), pCRFpoaeT534(2-2), and CLONE: pCRFpoaeT56(3-1)

FEATURE:
NAME/KEY: misc_feature
LCCATION: 1..30
OTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: /note= "5' end of large subunit OTHER INFORMATION: rRNA gene"
US-08-887-480-96
    INDIVIDUAL ISOLATE: T-427, T-534, and T-756 (INDIVIDUAL ISOLATE: sequence)
INMEDIATE SOURCE:
CLONE: DCRFpoaeT427(1-2), pCRFpoaeT534(2-2),
CLONE: pCRFpoaeT756(3-1)
                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: 181._337
CHER INFORMATION: /note= "5.85 rRNA gene"
FEATURE:
                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc feature
LOCATION: 31. 180
OTHER INFORMATION: /note= "ITS
                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
LOCATION: 338.7489
OTHER INFORMATION: /note= "ITS
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NAME/KEY: misc_feature
LOCATION: 490..546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 84.5
Matches 262; Conservative
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                                                                                                                                                                                                          318 Trececceccaciatricisesesesiscistresascerericaseses 377
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                                                                                    1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATCTTTGAACGCACA
                                                                                                           121 CC----CGGGTTTGGTGTTGGGGATCGGCAAG----CCCTTGCGGCAAGCCGGCCCCGAA
                                                                                                                                                                                                                                                                                                                                                                                                                  CGCGGCGCGCCGAAGCCGCTTAAACCCCCAACTTCTGAAT-GTTGACCTCGGATCAGGTAG
                                               Gaps
                                               11;
    DB 1; Length 594;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 96, Application US/08087480

Patent No. 5014453

GENERAL INFORMATION:
APPLICANT Beck, James J

TITLE OF INVENTION: Detection of Fungal Pathogens Using the TITLE OF INVENTION: Polymerase Chain Reaction
NUMBER OF SEQUENCES: 96

CORRESPONDENCES: 96

CORRESPONDENCES: No. 5814453artis Corporation
STREET: 520 White Plains Road

CITY: Tarrytown
                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,480
  Score 197.8; DB 1
Pred. No. 3.2e-56;
0; Mismatches 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08 08/722,187
FILING DATE: 15-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meige J. Timothy
REGISTRATION NUMBER: 38,241
REFFERENCE/DOCKET NUMBER: CGC 1739/PCT/CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    292 GAATACCCGCTGAACTTAA 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           556 GACTACCCGCTGAACTTAA 574
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LENGTH: 546 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLGY: linear
MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
STRAIN: FUBATIUM poae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
Similarity 85.0%;
1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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STATE: NY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10591
                                    271;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 9
US-08-887-480-96
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                                                                                                                                                                                                                                                                                          378
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    Query Match
                     Best Loca
Matches
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Gaps

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241 GGCCAAGCCGTTAAACCCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 300
        399 GCGGTCACG-TCGAGCTTCCATAGCGTAGTAATTTACACATCGTTACTGGTAATCGTCGC 457
                                                                   Sequence 82, Application US/08887480

Patent No. 5814453

GENERAL INFORMATION:

APPLICANT BECK, James J

TITLE OF INVENTION:

TITLE OF INVENTION:

NUMBER OF SEQUENCES: 96

CORRESPONDENCES: 96

CORRESPONDENCES: 36

CORRESPONDENCES: ADDRESSEE: No. 5814453artis Corporation

STATE: NY

CONTY: 1 AITYCOM

STATE: NY

COUNTY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 435
PIOR APPLICATION 1935
PIOR APPLICATION NUMBER: US 08/722,187
FILING DATE: 15-CCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timochy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: CGC 1739/PCT/CIP
TELEPHONE: 919-541-8587
TELEPHONE: 919-541-8689
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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LOCATION: 162.7318
OTHER INFORMATION: /note= "5.8S rRNA gene"
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INDIVIDUAL ISOLATE: R-5106, R-5126, and
INDIVIDUAL ISOLATE: (consensus seguence)
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LOCATION: 13..161
OTHER INFORMATION: /note= "ITS 1"
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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LOCATION: 1..12
OTHER INFORMATION: /note= "3'
OTHER INFORMATION: rRNA gene"
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
ORGANISM: Pusarium culmorum
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NAME/KEY:
LOCATION:
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US-08-887-480-82
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Best Local Similarity 84.5%; Pred. No. 8.1e-53;
Matches 262; Conservative 0; Mismatches 34; Indels 14; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Fusarium poae
INDIVIDUAL ISOLATE: T-427, T-534, and T-756 (consensus
INDIVIDUAL ISOLATE: sequence)
IMMEDIATE SOURCE:
CLONE: pCRFpoaeT427(1-2), pCRFpoaeT534(2-2), and
CLONE: pCRFpoaeT756(3-1)
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CCATION: 490..546
COTHER INFORMATION: /note= "5' end of large subunit;
CTHER INFORMATION: rRNA gene"
US-08-905-314A-22
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NAME/KEY: misc feature
LOCATION: 1.30
OTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
NAME/KEY: misc_feature
LOCATION: 181..337
OTHER INFORMATION: /note= "5.8S rRNA gene"
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NAME/KEY: misc feature
LOCATION: 31.180
OTHER INFORMATION: /note= "ITS 1"
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OTHER INFORMATION: /note= "ITS
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MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
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                                                                                                                                                                                    Gaps
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TITLE OF INVENTION: DETECTION OF WHEAT FUNGAL PATHOGENS
TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: 10. No. 5827653rtis Corporation Patent Department
STREET: 3054 Cormallis Road
CITY: Research Triangle Park
                                                                                                                                                                                    12;
                                                                                                                                                Length 504;
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                                   NAME/KEY: misc_feature
LOCATION: 473.504
LOCHER INFORMATION: Anote= "5' end of large subunit
OTHER INFORMATION: RNA gene"
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATR:
APPLICATION NUMBER: US/08/905,314A
                                                                                                                                             Score 186.6; DB 1;
Pred. No. 1.7e-52;
1; Mismatches 40;
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ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy
REGISTRATION UNUBER: 38,21
REFRENCE/DOCKET NUMBER: CGC 1944
TELECOMMUNICATION:
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   /note= "ITS
                                                                                                                                             Query Match
Best Local Similarity 82.6%;
Matches 252; Conservative
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TELEFAX: (919) 541-8689
INFORMATION FOR SEO ID NO: 19
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
OTHER INFORMATION: FEATURE:
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                                                                                                           US-08-887-480-82
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61 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC 120
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Sequence 82, Application US/08722187

Sequence 82, Application US/08722187

Patent No. 5955274

GREERAL INFORMATION:

APPLICANT: Ligon, James M

APPLICANT: Beck, James J

TITLE OF INVENTION: Detection of Fungal Pathogens Using the TITLE OF INVENTION: Detection of Fungal Pathogens Using the TITLE OF INVENTION: Polymerase Chain Reaction

NUMBER OF SEQUENCES: 86

CORRESPONDENCE ADDRESS:

GORRESPONDENCE ADDRESS:

STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATC
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                                                                                                                                                   FEATURE:
NAME/KEY: misc_feature
LOCATION: 1..12
COTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               end of large subunit
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Pred. No. 1.7e-52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: /note= "5.8S rRNA gene" FEATURE:
                                                                  ORGANISM: Fusarium culmorum INDIVIDUAL ISOLATE: R-5106, R-5126, and INDIVIDUAL ISOLATE: (consensus sequence)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Mismatches
                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: 13..161
OTHER INFORMATION: /note= "ITS 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LOCATION: 473. 504
; OTHER INFORMATION: /note= "5'
; OTHER INFORMATION: rRNA gene"
US-08-905-314A-19
                      MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
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82.6%;
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LOCATION: 473..504
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LOCATION: 319.7472
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Best Local Similarity
linear
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USA
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internal transcribed spacer region of Fusarium culmorum
(fculm.con)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            212 AAATGCGATAAGTAAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 186.6; DB 2; Length 504;
Pred. No. 1.7e-52;
1; Mismatches 40; Indels 12;
                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                   CLASSIFICATION: 3-3-3
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/233,608
FILING DATE: 04-APR-1994
ATTORNEY-AGENT INFORMATION:
NAME: WAlsh, Andrea C.
REGISTRATION NUMBER: 34,988
REFERENCE/DOCKET NUMBER: CGC 1739
TELECOMMUNICATION:
TELEPHONE: 919-541-8666
INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
LENGTH: 504 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 60.2%;
Best Local Similarity 82.6%;
Matches 252; Conservative
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LOCATION: 1..504

COTHER INFORMATION: interi

OTHER INFORMATION: interi

COTHER INFORMATION: (fcult

US-08-722-187-82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
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   Hawthorne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linea MOLECULE TYPE: DN HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTGAA 504
                                    USA
                                                     10532
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PCT-US95-04712-82
CITY: Har
STATE: N
COUNTRY:
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Sequence 82, Application PC/TUS9504712 GENERAL INFORMATION: APPLICANT: Ligon, James M

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/note= "DNA sequence for the
internal transcribed spacer region of Fusarium culmorum
(fculm.con)"
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APPLICANT: Beck, James J
TITLE OF INVENTION: Detection of Fungal Pathogens Using the
TITLE OF INVENTION: Polymerase Chain Reaction
TUMBER OF SEQUENCES: 86
CORRESPONDENCE 86
CORRESPONDENCE ADDRESS:
ADDRESSEE: Ciba-Geigy Corporation
STREET: 7 Skyline Drive
CITY: Hawthorne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 504;
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                                                                                                                                                                                                                                         ZIP: 10532
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40;
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                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FULUSSS/04/12
FULING DATE:
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: US 08/233,608
FILING DATE: 04-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Walsh, Andrea C.
REGISTRATION NUMBER: 34,988
REPERENCE/DOCKET NUMBER: GGC 1739
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8666
INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
LENGTH: 554 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature;
COCATION: 1..504
COTHER INFORMATION: inter:
CTHER INFORMATION: inter:
CTHER INFORMATION: (fculi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 504 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity 82.6'
Matches 252; Conservative
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283 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 342
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                                                              Sequence 4, Application US/08986727

Sequence 1, Application US/08986727

GENERAL INFORMATION:
APPLICANT: BROEL, Stacia R.
APPLICANT: IRELAM, Nancy A.
ITITLE OF INVENTION: DETECTION OF FUNGAL PATHOGENS
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS, L.L.P.
STREET: P.O. Box 1404
CITY: Alexandria
STATE: Victimia
STATE: Victimia
COUNTRY: United States
ZIP: 22313-1404
COMPUTER: IBM PC compatible
COMPUTER: IB
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MOLECULE TYPE: DNA (genomic)
RESULT 15
US-08-986-727-4
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Search completed: October 23, 2005, 16:07:06 Job time: 185.825 secs

121 CCCGGGTTTGGTGTTGGGGATCGGCAAGCCCTTGCGGCAAGCCGGCCCCCGAAATCTAGTG 180

578 CCGCTGAACTTAA 590

298 CCGCTGAACTTAA 310

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October 23, 2005, 15:20:46; Search time 1340.32 Seconds (without alignments) 1907.926 Million cell updates/sec
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1: /cgn2_6/ptodata/2/pubpna/USO7_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/USO7_BW PUB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9772231 segs, 4124568258 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                       OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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310
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Perfect score:
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                                                                                                                                                                                                                                                                      Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

			Description	e 7,		7	7	,	Sequence 5, Appli	Sequence 5, Appli
SUMMARIES			ΠD	US-10-046-955-7	US-09-961-755A-7	US-10-773-904-7	US-10-773-905-7	US-10-046-955-6	US-10-829-661-5	US-10-672-300-5
			DB	15	10	20	20	15	21	21
			Length	310	534	534	534	319	552	552
	dр	Query	re Match Length DB ID	100.0	7.76	7.76	7.76	77.5	75.3	75.3
			Score	310	302.8	302.8	302.8	240.2	233.4	233.4
		Result	No.		7	e	4	ഹ	9	7

) ORGANISM: Fusarium moniliforme US-10-046-955-7

LENGTH: 310

, 8 5, 13, 13, 13, 13, 13, 13, 13, 13, 13, 13	Sequence 6, Appli Sequence 6, Appli Sequence 1, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3, Appli Sequence 3, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 4, Appli Sequence 4, Appli Sequence 4, Appli Sequence 5, Appli Sequence 5, Appli Sequence 14, Appli Sequence 16, Appli Sequence 17, Appli Sequence 17, Appli Sequence 18, Appli Sequence 19, Appli Sequence 29, Appli Sequence 29, Appli Sequence 24, Appli Sequence 24, Appli Sequence 24, Appli Sequence 24, Appli Sequence 24, Appli Sequence 24, Appli
US-09-961-755A- US-09-961-755A- US-10-773-904-5 US-10-773-904-8 US-10-773-905-8 US-10-773-905-8 US-10-773-905-8	10 US-09-961-755A-6 20 US-09-961-755A-6 21 US-10-773-905-6 21 US-10-773-905-6 21 US-10-773-905-6 21 US-10-773-905-1 21 US-10-672-300-1 21 US-10-623-42-2 22 US-10-746-955-2 23 US-10-746-955-2 24 US-10-746-955-2 25 US-10-746-955-2 26 US-10-757-093-29 27 US-09-766-173C-4 28 US-09-766-173C-4 29 US-09-766-173C-5 20 US-09-766-173C-5 20 US-09-766-173C-5 20 US-09-961-663-14 20 US-09-961-663-15 20 US-09-961-663-15 20 US-09-961-663-15 20 US-09-961-663-17 21 US-10-046-955-25 21 US-09-961-663-17 21 US-10-046-955-25 21 US-09-961-663-17 21 US-10-046-955-25 22 US-09-961-663-17 23 US-09-961-663-17 24 US-09-961-663-17 25 US-09-961-663-17 26 US-09-961-663-17 27 US-09-961-663-17 28 US-09-961-663-17 28 US-09-961-663-17 28 US-09-961-663-17 28 US-09-961-663-17 28 US-09-961-663-17
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RESULT 1

US-10-046-955-7

Sequence 7, Application US/10046955

Publication No. US20030129600A1

GENERAL INPORMATION:

APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: Control and Prevention

APPLICANT: Morrison, Christine J.

APPLICANT: Morrison, Christine J.

APPLICANT: Morrison, Christine J.

APPLICANT: Addrevich, Liliana

APPLICANT: Ad

ALIGNMENTS

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APPLICANT: Barnett, Jason
TITLE DE INVENTION: Detection of Fusarium Species infecting Corn Using the TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/01/773,905
CURRENT FILING DATE: 2004-02-06
PRIOR PALICATION NUMBER: US/09/961,755
PRIOR FILING DATE: 2001-09-24
SUTHER FOR FILING DATE: 2001-09-24
SOFTWARE: Patent In version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 7, Application US/10773905; Publication No. US20040259121A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Fusarium proliferatum
US-10-773-904-7
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                                         529 CTGAAC 534
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APPLICANT: Beck, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/09/961,755A
CURRENT PILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
                                                                                                                        1 AAATGCGATAAGTAATGTGAATTGGAAAATTCAGTGAATCATCGAATCTTTGAACGCACA 60
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    Length 310;
                                         Indels
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; Score 310; DB 15;
; Pred. No. 2.1e-102;
0; Mismatches 0;
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Pred. No. 1.1e-99;
0; Mismatches 2;
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    100.0%;
100.0%;
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Best Local Similarity 99.3%;
Matches 304; Conservative
Query Match
Best Local Similarity 100.*
Matches 310; Conservative
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US-09-961-755A-7
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US-09-961-755A-7
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Sequence Journal of Application US/10773904
; Bublication No. US20040259120A1
; Berneral INFORMATION:
; GENERAL INFORMATION:
; APPLICANT: Beck, Jim
; APPLICANT: Barnett, Jason
; TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
; TITLE OF INVENTION: Polymerase Chain Reaction
; TITLE OF INVENTION: Polymerase Chain Reaction
; TITLE OF INVENTION: Polymerase Chain Reaction
; FILE REFERENCE: 60055
; CURRENT APPLICATION NUMBER: US/09/961,755
; PRIOR APPLICATION NUMBER: US/09/961,755
; PRIOR FILING DATE: 2004-02-06
; RAIDR FILING DATE: 2001-09-24
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin version 3.0
; FEG ID NO 7
; FEM OF THE OF
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                                                 63 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTACAACCCTCAGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 5, Application US/10829661
; Bublication No. US2005000951A1
; Bublication No. US2005000951A1
; GENERAL INFORMATION:
; APPLICANT: HAN, XIANG-YANG
; APPLICANT: TARRAND, JEFFREY J.
APPLICANT: PHAM, AUDREY S.
; APPLICANT: PHAM, AUDREY S.
; APPLICANT: PHAM, AUDREY S.
; TITLE OF INVENTION: DIAGNOSIS OF MOULD INFECTION
FILE REFERENCE: UTSC865US
; CURRENT APPLICATION NUMBER: US/10/829,661
; CURRENT APPLICATION NUMBER: 10/672,300
PRIOR FILING DATE: 2003-09-26
; PRIOR FILING DATE: 2003-09-26
; PRIOR FILING DATE: 2003-09-27
; NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                                                                         294 ATACCCGCTGAACTTAA 310
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SEQ ID NO 5
LENGTH: 552
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; ORGANISM: Fusarium sp.
US-10-829-661-5
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US-10-829-661-5
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APPLICANT: Control and Prevention.

APPLICANT: Morrison, Christine J.

APPLICANT: Adorevich, Liliana
APPLICANT: Aidorevich, Liliana
APPLICANTON NUMBER: US/10/046,955
CURRENT APPLICATION NUMBER: US/0423,233
FRIOR FILING DATE: 2000-06-27
PRIOR FILING DATE: 1998-05-01
PRIOR FILING DATE: 1998-05-01
PRIOR FILING DATE: 1998-05-01
PRIOR FILING DATE: 1997-06-02
NUMBER OF SEQ ID NOS: 61
SEQ ID NO 6
SEQ ID NO 6
LENGTH: 319
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97.7%; Score 302.8; DB 20;
Best Local Similarity 99.3%; Pred. No. 1.1e-99;
Matches 304; Conservative 0; Mismatches 2;
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Pred. No. 7.2e-77;
0; Mismatches 33
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Publication No. US20030129600A1
GENERAL INFORMATION:
                                          TYPE: DNA ORGANISM: Fusarium proliferatum
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Best Local Similarity 87.4%;
Matches 277; Conservative
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Publication No. US20030113722A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Beart, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the TITLE OF INVENTION: Polymerase Chain Reaction
TITLE OF INVENTION: Polymerase Chain Reaction
TITLE OF INVENTION: Polymerase Chain Reaction
GURRENT APPLICATION NUMBER: US/09/961,755A
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
SEQ ID NO 5
LENGTH: 522
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 552;
                                                                                Sequence 5, Application US/10672300; Bublication No. US20050048509A1; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: HAN, XIANG-YANG; APPLICANT: HAN, XIANG-YANG; TITLE OF INVENTION: DIAGNOSIS OF INVASIVE MOLD INFECTION FILE REFERENCE: UTXC:766US; CURRENT APPLICATION NUMBER: US/10/672,300; CURRENT FILING DATE: 2003-09-26; PRIOR APPLICATION NUMBER: 60/414,008; PRIOR PLING DATE: 2002-09-27; NUMBER OF SEQ ID NOS: 6; SOFTWARE: PATENTIN Ver: 2.1
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 536 ATACCCGCTGAACTTAA 552
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Matches 279; Conservative
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; ORGANISM: Fusarium sp.
US-10-672-300-5
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APPLICANT: Beck, Jim
APPLICANT: Beck, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/09/961,755A
CURRENT PILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
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DB 10; Length 522;
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US-09-961-755A-8
      Score 194.4; DB 10;
Pred. No. 5.1e-60;
0; Mismatches 36;
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Query Match 62.7%;
Best Local Similarity 84.3%;
Matches 258; Conservative
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ORGANISM: Fusarium subglutinans US-10-773-905-5
                                                                             Query Match 62.7%;
Best Local Similarity 84.3%;
Matches 258; Conservative
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                 ; TYPE: DNA
; ORGANISM: Fu
US-10-773-904-8
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LENGTH: 522
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Bernett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Detection of Fusarium Species infecting the
TITLE OF INVENTION: Polymerase Chain Reaction
TITLE OF INVENTION: Polymerase Chain Reaction
CURRENT APPLICATION WUMBER: US/10/773,904
CURRENT APPLICATION WUMBER: US/09/961,755
FRIOR APPLICATION WUMBER: US/09/961,755
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
SEQ ID NO 8
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APPLICANT: Beck, Jim
APPLICANT: Beck, Jim
APPLICANT: Beck, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Detection
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/10/773,904
CURRENT FILING DATE: 2004-02-06
PRIOR APPLICATION NUMBER: US/09/961,755
PRIOR PLILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn version 3.0
SEQ ID NO 5
LENGTH: 522
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC 120
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Pred. No. 5.1e-60;
0; Mismatches 36
                                                                                                            Sequence 5, Application US/10773904 Publication No. US20040259120A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Fusarium subglutinans
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Best Local Similarity 84.3%;
Matches 258; Conservative
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CTGAAC 306
                             CTGAAC 522
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US-10-773-904-5
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; Sequence 5, Application US/10773905; Sequence 5, Application US/10773905; Sequence 5, Application No. US20040259121A1
; GENERAL INFORMATION:
; APPLICANT: Back, Jim
; APPLICANT: Back, Jim
; TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
; TITLE OF INVENTION: Polymerase Chain Reaction
; FILE REFERENCE: 60055
; CURRENT APPLICATION NUMBER: US/10/773,905
; CURRENT FILING DATE: 2004-02-06
; PRIOR FILING DATE: 2001-09-24
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 5.2 348 120 121 CCCGGGTTTGGTGTTGGGGATCGCCAAGCCCTTGCGGCAAGCCGGCCCCCGAAATCTAGTG 180 GCGGTCTCGCTGCAGCTTCCATTGCGTAGTAATAAAACCCTCGCAACTGGTACGCGGCGC 240 457 GGCCAAGCCGTTAAACCCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 300 GGCCACGCCGTTAAA - CCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 516 121 CCCGGGTTTGGTGTTGGGGATCGCCAAGCCCTTGCGGCAAGCCGGGCCCCGAAATCTAGTG 180 181 GCGGTCTCGCTGCAGCTTCCATTGCGTAGTAAAACCCTCGCAACTGGTACGCGGCGC 240 229 AAATGCGATAAGTAAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 289 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC GCGGTCACG-TCGAGCTTCCATAGCGTAGTAAAACCCTCGTTACTGGTAATCGTCGC TTGCGCCCGCCAGTATTCTGGCGGCATGCCTGTTCGAGCGTCATTTCAACCCTCAAGCC 229 AAATGCGATAAGTAATGTGAATTGCAGAATTCAGTGAATCATCGAATCTTTGAACGCACA 289 TTGCGCCCGCCAGTATTCTGGCGGGCATGCCTGTTCGAGCGTCATTCAACCCTCAAGCC 1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATCTTTGAACGCACA 1 AAATGCGATAAGTAATGTGAATTGCAAAATTCAGTGAATCATCGAATCTTTGAACGCACA Gaps Gaps 12; 12; Query Match 62.7%; Score 194.4; DB 20; Length 522; Best Local Similarity 84.3%; Pred. No. 5.1e-60; Matches 258; Conservative 0; Mismatches 36; Indels 12; Length 522; Score 194.4; DB 20; Length Pred. No. 5.1e-60; 0; Mismatches 36; Indels Fusarium verticillioides (syn. F. moniliforme)

Gaps

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721 124

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229 AAATGCGATAAGTAATGGAGAATTCAGGGAATCATCGAATCTTGAACGCACA 288
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Publication No. US20030113722A1

GENERAL INFORMATION:

APPLICANT: Back, Jim

APPLICANT: Barnett, Jason

TITLE OF INVENTION: Polymerase Chain Reaction

FILE REFRENCE: 60055

CURRENT APPLICATION NUMBER: US/09/961,755A

NUMBER OF SEQ ID NOS: 24
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                                                                                                                                                                                                                                                                                                                                                                     Score 181; DB 20; Length 780;
Pred. No. 4.8e-55;
0; Mismatches 56; Indels
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Pred. No. 1.8e-54;
0; Mismatches 38;
                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Clone ID: MRT4577_57087C.1
US-10-425-115-137730
                                                                                                                                                                                                                                            LOCATION: (1)...(780)
OTHER INFORMATION: unsure at all n locations
  TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 137730
LENGTH: 780
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SEQ ID NO 6
LENGTH: 521
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Best Local Similarity 79.1%;
Matches 242; Conservative (
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Best Local Similarity 83.3%;
Matches 255; Conservative
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                                                                                                                                                        TYPE: DNA
ORGANISM: Zea mays
                                                                                                                                                                                                                          NAME/KEY: unsure
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; GENERAL INFORMATION:
; APPLICANT: Beck, Jim
; APPLICANT: Barnett, Jason
; TITLE OF INVENTION: Polymerase Chain Reaction
; TITLE OF INVENTION: Polymerase Chain Reaction
; FILE REFERENCE: 60055
; CURRENT APPLICATION NUMBER: US/10/773,905
; FILE REFERENCE: 2004-02-06
; PRIOR FILING DATE: 2004-02-06
; PRIOR FILING DATE: 2001-09-24
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 8
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US-10-425-115-137730/c
; Sequence 13773, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: La Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              240
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GCCCAAGCCGTTAAACCCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 300
                                                                                                           GGCCACGCCGTTAAA-CCCCCAACTTCTGAATGTTGACCTCGGATCAGGTAGGAATACCCG 516
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ORGANISM: Fusarium verticillioides (syn. F. moniliforme)
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Search completed: October 23, 2005, 23:48:15 Job time : 1340.32 secs

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; TYPE: DNA
; ORGANISM: Fusarium moniliforme
US-09-423-233-7
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Best Local Similarity 94.4
Matches 17; Conservative
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US-09-423-233-7
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Sequence 8, Appli
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Sequence 1326, Appli
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Sequence 3309, Ap
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Sequence 2042, Ap
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3055, Ap
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3385, Ap
4248, Ap
16629, A
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                                                         October 23, 2005, 06:56:31 ; Search time 10.7318 Seconds (without alignments) 2744.464 Million cell updates/sec
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/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
       GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-08-652-127C-5
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US-08-652-127C-6
US-08-652-127C-6
US-09-949-016-13246
US-09-949-016-17536
US-09-949-016-17536
US-09-949-016-54231
US-09-170-279-3055
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US-09-170-769A-5
US-08-737-129A-5
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PCT-US94-14106-50
PCT-US94-14106-54
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US-08-737-129A-1
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Maximum Match 100%
Listing first 45 summaries
                                        - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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Match Length DB
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30, Appl
1514, Ap
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Patent No. 6372430
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL TOWS:
TITLE OF INVENTION:
TITLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 03063-0341WP
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 61

SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/09423233
Sequence 7, Application US/09423233
Patent No. 6372430
GENERAL INFORMATION
GENERAL INFORMATION
APPLICANT: The Government of the United States of America as
TITLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 03063-0341WP
CURRENT APPLICATION NUMBER: US/09/423,233
NUMBER OF SEQ ID NOS: 2000-06-27
SOFTWARE: PatentIn Ver. 2.0
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US-09-011-769A-20
US-08-133-011-99
US-08-37-87A-99
US-08-387-619-99
US-08-907-739-99
US-09-729-597-99
PCT-US93-08364-71
US-08-133-011-115
US-08-133-011-115
US-08-132-730A-115
US-08-387-619-115
US-08-387-619-115
US-08-907-739-115
US-08-907-739-115
US-09-729-597-115
US-09-729-597-115
US-09-739-08364-88
PCT-US93-08364-88
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US-10-134-188-30
US-10-134-188-30
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Score 16.4; DE Pred. No. 17; 0; Mismatches

91.1%; 94.4%;

Length 310,

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Sequence 7, Application US/08652127C; Patent No. 5792611; GENERAL INFORMATION: APPLICANT: Richard C. Hamelin TITLE OF INVENTION: PATHOGEN FUNGI (CORRESPONDENCE LADRESSE: ADDRESSE: Seaby & MacLean; ADDRESSE: Seaby & MacLean
                                  OPERATING SISTEL.
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: George A. Seaby
REGISCHARTION NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
TELEPHONE: (613) 232-5815
TELEPRAX: (613) 232-5815
IBM PC Compatible
SYSTEM: PC-DOS/MS-DOS
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ORGANISM: Gibberella fujikuroi
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Best Local Similarity 94.4%;
Matches 17; Conservative
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STRANDEDNESS: double
   COMPUTER: IBM PC OPERATING SYSTEM:
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SEQ ID NO 1
LENGTH: 2293
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US-08-652-127C-5
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                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: George A. Seaby
ADDRESSEE: George A. Seaby
ADDRESSEE: George A. Seaby
ADDRESSEE: George A. Seaby
CITY: Octawa
CITY: Octawa
CITY: Crawa
ZIP: KIR 6K7
COMPUTRY: Canada
ZIP: KIR 6K7
COMPUTRY: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: LEM PC Compatible
COMPUTER: ACTIVE
COMPUTER: ACTIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 4

US-08-652-127C-5

Sequence 5, Application US/08652127C

Sequence 5, Papplication US/08652127C

Patent No. 5/92611

GENERAL INFORMATION:

APPLICANT: Richard C. Hamelin

TITLE OF INVENTION: DETECTION OF PLANT

TITLE OF INVENTION: PATHOGEN FUNGI

NUMBER OF SEQUENCES: 10

CORRESPONDENCE ADDRESS:

ADDRESSEE: George A. Seaby

ADDRESSEE: Seaby & MacLean

STREET: 880 Wellington Street, Suite 708

CITY: Octawa

COUNTRY: Canada

ZIP: KIR 6K7

COUNTRY: READABLE FORM:

MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91.1%; Score 16.4; 94.4%; Pred. No. 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: George A. Seaby
REGISCHARTION NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 1898
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
TELEPRAX: (613) 232-5815
TELEFAX: (613) 232-5815
TELEFAX: (613) 232-5815
TELEFAX: (513) 232-5815
                                                                                                                                                                                                                                                    US-08-652-127C-8
; Sequence 8, Application US/08652127C
; Patent No. 5792611
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                                                                                                       174 rczacrececierczeci 191
                                      1 TCTAGTGACGGTCTCGCT 18
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Best Local Similarity 94.4
Matches 17; Conservative
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STRANDEDNESS: double
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Sequence 1, Application US/09645073

Fatent No. 6287800

FREENRAL INFORMATION:
APPLICANT: Lee, May
APPLICANT: Lee, May
APPLICANT: Leo, May
CURRENT APPLICANT: D.2-01NP
CURRENT APPLICATION NUMBER: US/09/645,073

CURRENT FILING DATE: 2000-08-25

PRIOR PELICATION NUMBER: US 60/151,770

PRIOR FILING DATE: 1999-08-31

NUMBER OF SEQ ID NOS: 1
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Score 16.4; DB 1; Length 594;
Pred. No. 18;
0; Mismatches 1; Indels
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US-09-949-016-13246

i Sequence 13246

patent No. 681239

i Patent No. 681239

i GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

ITILE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

ITILE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

ITILE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT PILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/231,768

PRIOR PILING DATE: 2000-10-03

PRIOR PILING DATE: 2000-10-03

PRIOR PILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FasteEQ for Windows Version 4.0

SEQ ID NO 13246
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US-09-49-016-17536/c

US-09-494-016-17536, Application US/09949016

Sequence 17536, Application US/09949016

Sequence 17536, Application US/09949016

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR PILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                 Length 581;
                                                                                                                                                 2; Indels
                                                                                               Score 14.8; DB 1;
Pred. No. 1.4e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                             427 TATAGTGGCGGTCTCGCT 444
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                                                                                                 82.2%;
                                                                                            Query Match
Best Local Similarity 88.9°
Matches 16, Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
       double
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NAME/KEY: misc_feature
LOCATION: (1)...(39376)
  ; STRANDEDNESS: dou
; TOPOLOGY: linear
US-08-652-127C-6
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ORGANISM: Human
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ORGANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6, Application US/08652127C

Sequence 7, Application US/08652127C

Patent No. 5792611

SENERAL INFORMATION:
APPLICANT: Richard C. Hamelin
TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: PATHOGEN FUNGI
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seaby & MacLean
STREET: 880 Wellington Street, Suite 708
CITY: Ottawa
COUNTY: Canada
ZIP: KIR 6K7
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION NUMBER: 1998
TELECOMMUNICATION INFORMATION:
TELEBRAX: (613) 232-5815
TELEFRAX: (613) 23-5831
INFORMATION FOR SEQ ID NO: 6:
880 Wellington Street, Suite 708
       CITY: Ottawa
CITY: Ottawa
COUNTRY: Canada
ZIP: KIR 6K7
COMPUTER: BADABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,127C
FILING DATE: May 23, 1996
CLASSIFICATION 1435
ATTONEY/AGENT INFORMATION:
NAME: George A. Seaby
REGISTRATION NUMBER: 24,034
REGISTRATION NUMBER: 1898
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              377 TATAGTGGCGGTCTCGCT 394
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STRANDEDNESS: double
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US-08-652-127C-7
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Gaps

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Indels

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1; Mismatches

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15; Conservative
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US-09-252-991A-2042
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       Matches
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i Sequence 5431, Application US/09949016

j Fatent No. 681239

j GENERAL INFORMATION:

i APPLICANT: VENTER,

i TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

i TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

i TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

i CURRENT APPLICATION NUMBER: US/09/949,016

i CURRENT FILING DATE: 2000-04-14

i PRIOR PAPLICATION NUMBER: 60/231,768

i PRIOR FILING DATE: 2000-10-20

i PRIOR FILING DATE: 2000-09-08

i PRIOR FILING DATE: 2000-09-08

i NUMBER OF SEQ ID NOS: 207012

i SOFTHARE: FeatSEQ for Windows Version 4.0

i LENGTH: 601
                                                                                                                                                                                                                                                                                                              ; Sequence 16769, Application US/09949016
; Patent No. 6812339
; GENERAL INCORMATION:
    APPLICANT: VENTER, J. Craig et al.
    APPLICANT: VENTER, J. Craig et al.
    TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
    TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
    PILE REPERENCE: CL001307
    CURRENT APPLICATION NUMBER: US/09/949,016
    PRIOR APPLICATION NUMBER: 60/241,755
    PRIOR PLILING DATE: 2000-10-20
    PRIOR PLLING DATE: 2000-10-03
    PRIOR FILING DATE: 2000-10-03
    PRIOR FILING DATE: 2000-09-08
    NUMBER OF SEQ ID NOS: 207012
    SOFTWARE: FastSEQ for Windows Version 4.0
    SEQ ID NO 16769
    LENGTH: 63982
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                                                                      Score 14.8; DB 4; Length 39376;
Pred. No. 2.1e+02;
0; Mismatches 2; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     82.2%; Score 14.8; DB 4; Length 63982; 88.9%; Pred. No. 2.2e+02; rive 0; Mismatches 2; Indels 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19046 TCTAGTGACTGTCTCACT 19063
; OTHER INFORMATION: n = A, T, C \text{ or } G
US-09-949-016-17536
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                                                                      Query Match 82.2%;
Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
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Best Local Similarity
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ORGANISM: Human
                                                                                                                                                                                                                                                                                                         US-09-949-016-16769
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ORGANISM: Human
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PUB. 199480012
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SEG ID NO 3309
LENGTH: 945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: KIMMERLY, WILLIAM JOHN
APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT PILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SSEQ ID NO 3055
LENGTH: 1146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 80.0%; Score 14.4; DB 4; Length 945; Best Local Similarity 93.8%; Pred. No. 2.4e+02; Matches 15; Conservative 0; Mismatches 1; Indels
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                                                                                           US-09-710-279-3309/c
Sequence 3309, Application US/09710279
; Patent No. 6703492
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-710-279-3055/c
; Sequence 3055, Application US/09710279
; Patent No. 6703492
314 TTTÁGAGÁCGGTCYCGCT 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         268 CTACTGACGGTCTCGC 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           469 CTACTGACGGTCTCGC 454
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity 93.0%
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RESULT 15

US-09-252-991A-1914/C

Sequence 1914, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:

APPLICATION:

TITLE OF INVENTION:

TITLE OF INVENTION:

TITLE OF INVENTION:

TITLE OF INVENTION:

ARRIGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT APPLICATION NUMBER: US/00/224,991A

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR PILING DATE: 1998-02-18

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 1914

LENGTH: 1908
                                         GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 2042
LENGTH: 1629
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Sequence 2042, Application US/09252991A Patent No. 6551795
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ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Pseudomonas aeruginosa
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Best Local Similarity 93.81
Matches 15; Conservative
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                    - nucleic search, using sw model
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October 23, 2005, 15:20:46; Search time 77.8251 Seconds (without alignments) 1907.926 Million cell updates/sec 1 tctagtgacggtctcgct 18 US-10-046-955-49 18 Title: Perfect score: OM nucleic Sequence: Run on:

IDENTITY NUC Gapop 10.0 , Gapext 1.0 Scoring table:

19544462 Total number of hits satisfying chosen parameters: 9772231 segs, 4124568258 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Published Applications NA:*

1: \cgn2 \(\) \cgn2 \(\) \cgn \(\)

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	Sequence 49, Appl	Sequence 7, Appli	Sequence 7, Appli	Sequence 7, Appli	Sequence 7, Appli	Seguence 131520,	Sequence 33508, A
ΙD	US-10-046-955-49	US-10-046-955-7	US-09-961-755A-7	US-10-773-904-7	US-10-773-905-7	US-10-425-115-131520	US-09-918-995-33508
DB	15	15	10	20	20	20	10
% Query Match Length DB ID	18	310	534	534	534	279	386
% Query Match	100.0	91.1	91.1	91.1	91.1	85.6	83.3
Score	18	16.4	16.4	16.4	16.4	15.4	15
Result No.	1	~	m	4	Ŋ	9	c 7

US-10-046-955-49

Sequence 34287, A Sequence 1024, Ap Sequence 24124, A Sequence 237, App Sequence 568242,	2782, 74582, 215234 215235 215234 215235	35.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0 3	~	126 200, 254, 277 400, 88009
09-918-995-34287 09-918-995-1024 -10-450-763-24124 110-684-422-237 111-036-317-568242	-10-357-930-2782 -10-437-963-74582 -10-027-632-215234 -10-027-632-215235 -10-027-632-215235	10-76/701-8208 10-425-115-42609 -09-925-065A-35831 -09-925-065A-37892 -09-925-065A-880442 10-437-963-82321 110-741-600-17944 10-719-993-6767	10-087-192-1024 10-633-423-9 10-427-7419 10-087-192-166 59-880-107-3814 10-775-169-338 10-775-149-12 10-756-149-12	10-425-115-128972 110-719-900-176089 110-516-429-20 99-864-761-25477 110-276-774-20 09-925-065A-40631 09-925-065A-406632 99-864-761-8809
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ALIGNMENTS

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Sequence 49, Application US/10046955

Sequence 49, Application US/10046955

Publication No. US2030129600A1

GENERAL INFORMATION:

APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: Control and Prevention

APPLICANT: Control and Prevention

APPLICANT: Reiss, Errol

APPLICANT: Reiss, Errol

APPLICANT: Aidorevich, Liliana

APPLICANT: Aidorevi
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LENGTH: 18
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Sequence 7, Application US/10773904
Publication No. US20040259120A1
GENERAL INFORMATION:
APPLICANT: Beck, Jim
APPLICANT: Beck, Jim
APPLICANT: Becketion of Fusarium Species infecting Corn Using the TITLE OF INVENTION: Detection of Fusarium Reaction
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/10/773,904
CURRENT FILING DATE: 2004-02-06
PRIOR FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
NUMBER OF SEQ ID NOS: 24
NUMBER PETERIAL OF SEQ ID NOS: 24
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TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
FILE REFERENCE: 60055
CURRENT PELLON NUMBER: US/09/961,735
PRIOR PILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
SEQ ID NO 7
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Pred. No. 83;
0; Mismatches 1;
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Best Local Similarity 94.4%; Pred. No. 83;
Matches 17; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 131520, Application US/10425115; Publication No. US20040214272A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7, Application US/10773905; Publication No. US20040259121A1; GENERAL INFORMATION:
                                                                            402 TCTAGTGGCGGTCTCGCT 419
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               18
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, ORGANISM: Fusarium proliferatum
US-10-773-904-7
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; ORGANISM: Fusarium proliferatum
US-10-773-905-7
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Best Local Similarity 94.4%;
Matches 17; Conservative (
               1 TCTAGTGACGGTCTCGCT
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; Sequence 7, Application US/20030129600A1
; Sequence 7, Application No. US20030129600A1
; Sequence 7, Application No. US20030129600A1
; GENERAL INFORMATION:
    APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: Secretary of the Department of Health and Human Services, Centers for Dis APPLICANT: Mortison, Christine J.
    APPLICANT: Aidorevich, Liliana
    APPLICANT: Aidorevich, Liliana

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APPLICANT: Beck, Jim
APPLICANT: Barnett, Jason
TITLE OF INVENTION: Detection of Fusarium Species infecting Corn Using the
TITLE OF INVENTION: Polymerase Chain Reaction
FILE REFERENCE: 60055
CURRENT APPLICATION NUMBER: US/09/961,755A
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.0
SEQ ID NO 7
LENGTH::534
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         91.1%; Score 16.4; DB 15; Length 310; 94.4%; Pred. No. 85;
            DB 15; Length 18; 13;
                                                                        0; Indels
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               Score 18;
Pred. No.
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         100.0%;
                                                                                                                                                              1 TCTAGTGACGGTCTCGCT 18
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ORGANISM: Fusarium proliferatum
US-09-961-755A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Fusarium moniliforme
US-10-046-955-7
                                                                                                                                     1 TCTAGTGACGGTCTCGCT
Ouery Match
Best Local Similarity 100./
Matches 18; Conservative
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Best Local Similarity 94.4
Matches 17; Conservative
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Best Local Similarity
Matches 17; Conserva
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US-09-961-755A-7
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; Sequence 1024, Application US/09918995
; Publication No. US20030073623A1
; GENERAL INFORMATION:
    APPLICANT: Hyseq, Inc.
    TITLE OF INVENTION: PROM VARIOUS cDNA LIBRARIES
; TITLE OF INVENTION: FROM VARIOUS cDNA LIBRARIES
; FILE REPERENCE: 20411-756
; CURRENT APPLICATION NUMBER: US/09/918,995
; CURRENT FILING DATE: 1999-01-20
; PRIOR APPLICATION NUMBER: US/09/235,076
; RICH REPERENCE: 1999-01-20
; NUMBER OF SEQ ID NOS: 38054
; SOFTWARE: FBSESEQ for Windows Version 3.0
; SEQ ID NO 1024
                                                                                                                                                                                                                                  Query Match 83.3%; Score 15; DB 10; Length 449; Best Local Similarity 100.0%; Pred. No. 4.7e+02; Matches 15; Conservative 0; Mismatches 0; Indels
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; Publication No. US20550196754A1
; GENERAL INFORMATION:
    APPLICANT: HYPEG, Inc
    ITLE OF INVENTION:
    TITLE OF INVENTION NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
    TILE REPERENCE: 790CIP3/US
    CURRENT APPLICATION NUMBER: US/10/450,763
    CURRENT PILING DATE: 2003-66-11
    PRIOR APPLICATION NUMBER: PCT/US01/08631
    PRIOR FILING DATE: 2000-03-30
    PRIOR FILING DATE: 2000-03-31
    PRIOR PILING DATE: 2000-03-31
    PRIOR PILING DATE: 2000-03-31
    PRIOR PILING DATE: 2000-08-23
; WINDRER OF SEQ ID NOS: 60736
; SOFTWARE: CUSTOM
    SEQ ID NOS: 60736
; SEQ ID NO 24124
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83.3%; Score 15; DB
Best Local Similarity 100.0%; Pred. No. 4.7
Matches 15; Conservative 0; Mismatches
                                                                              FEATURE:
| NAME/KEY: misc_feature
| LOCATION: [1]...(449)
| COTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; LOCATION: (1).T. (473)
; OTHER INFORMATION: n = A,T,C or G
US-09-918-995-1024
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                                    TYPE: DNA
ORGANISM: Homo sapiens
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             LENGTH: 449
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Town, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Dlants
FITLE OF INVENTION: Dlants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 131520
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; Publication No. US20030073623A1
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc.
; TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
; TILLE OF INVENTION: ROW VARIOUS cDNA LIBRARIES
; FILE REPERENCE: 20411-756
; CURRENT APPLICATION NUMBER: US/09/918,995
; CURRENT FILING DATE: 2001-07-30
; PRIOR FILING DATE: 2001-07-30
; PRIOR FILING DATE: 1099-01-20
; NUMBER OF SEQ ID NOS: 38054
; SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hyeeq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: PROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT PILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 33508
LENGTH: 386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 83.3%; Score 15; DB 10; Length 386; Best Local Similarity 100.0%; Pred. No. 4.8e+02; Matches 15; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 94.1%; Pred. No. 2.9e+02;
Matches 16; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Clone ID: MRT4577_51429C.1
US-10-425-115-131520
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Publication No. US20030073623A1
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-09-918-995-33508
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TYPE: DNA
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Publication No. US20040229233A1

GENERAL INFORMATION:

APPLICANT: ABURATAIN:

APPLICANT: ABURATAIN:

APPLICANT: YAMAMOTO, Shogo

TITLE OF INVENTION: Human housekeeping genes and human tissue-specific genes

FILE REFERENCE: 113991

CURRENT FILING DATE: 2002-10-15

PRIOR APPLICATION NUMBER: US 60/418,614

PRIOR FILING DATE: 2002-10-16

NUMBER OF SEQ ID NOS: 332

SOFTWARE: Patentin version 3.2

SEQ ID NO 237
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Publication No. US20050214823A1

GENERAL INFORMATION:

APPLICANT: Williams, Alan

TITLE OF INVENTION: Method of Analysis of Alternative Splicing in Mouse

FILE REFERENCE: 3654.1

CURRENT APPLICATION NUMBER: US/11/036,317

CURRENT FILING DATE: 2005-01-13

PRIOR PILING DATE: 2004-01-13

PRIOR FILING DATE: 2004-01-13

NUMBER OF SEQ ID NOS: 991174

SSOFTWARE: Microarray Probe Sequence Listing Generator V 1.1

LENGTH: 25
                                                         LOCATION: (1416)..(1033)
COTHER INFORMATION: 64% homologous to Homo sapiens HCG V, accession number:
COTHER INFORMATION: U53588, Smith-Waterman Score=380.
US-10-450-763-24124
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100.0%; Pred. No. 4.6e+02;
tive 0; Mismatches 0; Indels C
                                                                                                                                                                   Length 1596;
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                                                                                                                                                              83.3%; Score 15; DB 22; I
100.0%; Pred. No. 4.6e+02;
iive 0; Mismatches 0;
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88.9%;
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                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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ORGANISM: Mus musculus
US-11-036-317-568242
ORGANISM: Homo sapiens
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; LOCATION: (200)..(580)
US-10-684-422-237
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Best Local Similarity
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US-11-036-317-568242/c
                   FEATURE:
NAME/KEY: SIMILAR
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US-10-684-422-237/c
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APPLICANT: La Rosa Thomas J
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Abu Yihua
APPLICANT: Abu Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223) B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
LENGTH: 388
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US-10-379-391-2782/C

US-10-379-3791-2782/C

US-10-379-3791-2782/C

Publication No. US20040259086A1

GENERAL INFORMATION:

APPLICANT: Endegel. Robert

APPLICANT: Endegel. Robert

APPLICANT: Mondan, John

TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR

TITLE OF INVENTION: HUMAN PROSTATE CANCER

FILE REPERENCE: MRI-007BCN

CURRENT APPLICATION WUMBER: 09/785,276

PRIOR APPLICATION NUMBER: 09/785,276

PRIOR APPLICATION NUMBER: 09/183,319

PRIOR APPLICATION NUMBER: 60/183,319

PRIOR FILING DATE: 2000-02-16

PRIOR PELING DATE: 2000-03-16

PRIOR PELING DATE: 2000-03-16

PRIOR PILING DATE: 2000-05-16

PRIOR PILING DATE: 2000-05-18

PRIOR PILING DATE: 2000-01-18

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US-10-424-599-44516
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    5,
    Mismatches
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; Sequence 44516, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
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    16; Conservative
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ORGANISM: Homo sapiens
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US-10-43/-963-4936/C
US-10-43/-963-4936/C
Sequence 7582, Application US/10437963
Fublication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Wu, Wei
APPLICANT: Barbazuk, Brad
APPLICANT: Bing
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 74582
LENGTH: 558
TYPE: DNA
PERMINE:
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                                                                                                                                Length 403;
                                                                                                                             Query Match
Best Local Similarity 88.9%; Pred. No. 6.1e+02;
Matches 16; Conservative 0; Mismatches 2; Indels
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US-10-437-963-74582
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; FEATURE:

; NAME/KEY: misc_feature

; LOCATION: 211, 216, 362, 364

; OTHER INFORMATION: n = A,T,C or G

US-10-357-930-2782
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US-10-437-963-74582/c
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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/PcTuS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
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1 cgttaattcgcgttcctc 18
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

st Query Score Match Length DB ID	E	E		SUMMAKI ES	Description
100.0 18 3	1	1)-SII	US-09-423-233-50	Semience 50. Anni
83.3 615 4	4	4 US-	SD-	US-09-134-000C-314	314
82.2 207 4	4	4 US	SD	US-09-107-532A-1902	1902
14.8 82.2 650 4 US	4	4 US	S	US-09-669-751-103	103, 7
4	4	4 UE	5	US-09-134-000C-2950	2950,
80.0 939 4	4	4 US	ă	US-09-710-279-3059	
14.4 80.0 963 3 US	<u>"</u>	3 US	S	US-09-134-001C-1367	Sequence 1367, Ap
14.4 80.0 2815 4 US	4	4 US	ns	US-09-710-279-3819	
.4 80.0 146401 4	4	4 US	ns	US-09-949-016-16151	Sequence 16151, A
-	-	1 US	ns	US-07-972-791-3	3,
14 77.8 3345 1 US-(-1	1 US	S	-07-972-791-7	7
14 77.8 3346 1 US	-	1 US	ns	US-07-972-791-5	'n
3347 1	П	1 US	ns	US-07-972-791-2	N
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3361 1	-	1 US	Ŕ	US-07-972-791-6	9
3434 6	9	6 53	53	5310649-1	٠.
77.8 3434 6	9	6 53	53	5310649-1	Patent No. 5310649
76.7 601 4	4	4 US	ns	US-09-949-016-142005	Seguence 142005,
76.7 601 4	4	4 US	ns	US-09-949-016-142006	Sequence 142006,
76.7 601 4	4	4 US	S	US-09-949-016-172751	Sequence 172751,
76.7 702 4	4	4 US	5	US-09-543-681A-4125	Seguence 4125, Ap
76.7 1356 4	4	4 US	S	US-09-328-352-1886	Seguence 1886, Ap
8 76.7 1386 3	m	3 US	S	US-09-134-001C-984	984, 7
76.7 2229 4	4	4 U	Ď	US-09-902-540-4193	4193
8 76.7 5679 3	m	3	ŏ	US-08-844-274-10	10, A
76.7 5679 4	4	4 US	SD	US-09-598-421-10	10,
76.7 6642 4 1	4	4 US	S	US-09-809-665A-80	Sequence 80, Appl

Sequence 13, Appl Sequence 14, Appl Sequence 14, Appl Sequence 20, Appl Sequence 20, Appl Sequence 6, Appl Sequence 1629, A Sequence 1, Appl Sequence 1, Appl Sequence 11862, A Sequence 1295, A	erica as rgillus Species and	Length 18; Indels 0; Gaps 0;	ENCES RELATING TO OSTICS AND THERAPEUTICS	Length 615;
US-08-844-274-13 US-08-844-274-14 US-09-598-421-13 US-09-598-421-14 US-08-844-274-20 US-09-598-421-20 US-09-377-066-6 US-09-377-066-6 US-09-377-066-6 US-09-949-016-16629 US-09-949-016-16862 US-09-949-016-12953 US-09-949-016-12954	ALIGNMENTS 9423233 the United States of America as wids for Detecting Aspergillus Lamentous Fungi SS/09/423,233	Score 18; DB 3; Pred. No. 1.9; 0; Mismatches 0; 18	num et al ACID AND AMINO ACID SEQUENCES I ACID AND AMINO ACID SEQUENCES I SECOUS FAECALIS FOR DIAGNOSTICS 13-13 60/055,778	Score 15; DB 4;
3.8 76.7 6723 3 3.8 76.7 6723 4 3.8 76.7 6723 4 3.8 76.7 750 3 3.8 76.7 7560 3 3.8 76.7 9560 4 3.8 76.7 970 3 3.8 76.7 970 3 3.8 76.7 970 4 3.8 76.7 46492 4 3.8 76.7 61847 4	RESULT 1 US-09-423-233-50 Sequence 50, Application US/09423233 Patent No. 6372430 GENERAL INFORMATION: TITLE OF INVENTION: Nucleic Acids for Deter TITLE OF INVENTION: Other Filamentous Fung; FILE REFERENCE: 03063-0341WP CURRENT APPLICATION NUMBER: US/09/423,233 CURRENT APPLICATION NUMBER: US/09/423,233 CURRENT APPLICATION NUMBER: US/09/423,233 SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 50 LANGTH: 18 LANGTH: 18 LANGTH: 18 LANGTH: DNA ORGANISM: Fusarium oxysporum US-09-423-233-50	Match 100.0%; Local Similarity 100.0%; les 18; Conservative 1 CGTTAATTCGCGTTCCTC 1 CGTTAATTCGCGTTCCTC 1 CGTTAATTCGCGTTCCTC 1 CGTTAATTCGCGTTCCTC	RESULT 2 US-09-134-000C-314 Sequence 314, Application US/09134000C Sequence 314, Application US/09134000C Patent No. 6617156 GENERAL INFORMATION: APPLICANT: Lynn Doucette-Stamm et al TITLE OF INVENTION: BUTERCOCCUS FAECALIS FOI FILE REPRENCE: 032796-032 CURRENT APPLICATION NUMBER: US/09/134,000C CURRENT APPLICATION NUMBER: US/09/134,000C CURRENT APPLICATION NUMBER: US/09/134,000C CURRENT FILING DATE: 1997-08-15 NUMBER OF SEQ ID NOS: 6812 SEQ ID NO 314 LENGTH: 615 SEQ ID NO 314 LENGTH: 615 TYPE: DAY ORGANISM: Enterococcus faecalis US-09-134-000C-314	th 83.3%;
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	RESULT 1 US-09-423-233- ; Sequence 50, ; Patent No. 6 GENERAL INFO ; TITLE OF	Query Match Best Local Matches 1 Qy 1	RESULT 2 US-09-134-000C-314, Ap ; Sequence 314, Ap ; Patent No. 66171 ; GENERAL INFORMAT ; TITLE OF INVENT ; FILM ; RICHARD DA ; SOFTWARE: Paten ; SEQ ID NO 314 ; LENGTH: 615 ; TYPE: DNA ; CRGANISM: Ente	Query Match

us-10-046-955-50.rni

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Query Match 82.2%;
Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
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US-09-669-751-103
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                                                                                                                                                                                  Sequence 1902, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                     Gaps
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Pred. No. 1.1e+02;
; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
CITY: Waltham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8; DB 4;
Pred. No. 1.3e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/05151
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) NAME/KEY: misc_feature
) LOCATION: (B) LOCATION 1...207
) SEQUENCE DESCRIPTION: SEQ ID NO: 1902:
US-09-107-532A-1902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: PC
OPERATING SYSTEM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: circular
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (781) 893-8277
INFORMATION FOR SEQ ID NO: 1902:
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
STRANDEDNESS: double
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COUNTRY: USA
ZIP: 02354
                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 7310
 100.08;
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                                                                            172 GTTAATTCGCGTTCC 186
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                                                      2 GTTAATTCGCGTTCC 16
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Best Local Similarity 88.9
Matches 16; Conservative
                 15; Conservative
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Best Local Similarity
                                                                                                                                                   RESULT 3
US-09-107-532A-1902/c
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                 Matches
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RESULT 4 US-09-669-751-103 ; Sequence 103, Application US/09669751

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84 cGrraarrcrcrrrccrc

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RESULT 5
US-09-114-000C-2950/C
i Sequence 2950, Application US/09134000C
j Patent No. 6617156
j GENERAL INPORMATION:
APPLICATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUMBER: US/09/134,000C
CURRENT FILING DATE: 1998-08-13
PRIOR FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 6812
SEQ ID NO 2950
LENGTH: 2457
LENGTH: 2457
Patent No. 6551575
GENBRAL INFORMATION:
APPLICANT: Greenspan, Ralph J.
TITLE OF INVENTION: Methods for Identifying Compounds for
TITLE OF INVENTION: Motion Sickness, Vertigo and Other Disorders Related to
TITLE OF INVENTION: Balance and the Perception of Gravity
TITLE OF INVENTION: Balance and the Perception of Gravity
FILE REFERENCE: P-NI 3864
CURRENT APPLICATION NUMBER: US 60969,751
CURRENT FILING DATE: 1999-12-02
NUMBER OF SEQ ID NOS: 261
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 103
LENGTH: 650
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US-09-110-279-3059/C
US-09-110-279-3059, Application US/09710279
Patent No. 6703492
GENERAL INFORMATION:
TATLE OF INVENTION: WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 1099-11-09
PRIOR PILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFWARE: Patentin Ver. 2.1
SEQ ID NO 3059
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88.9%; Pred. No. 1.6e+02;
tive 0; Mismatches 2; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 650;
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Pred. No. 1.4e+02;
0; Mismatches 2;
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US-09-134-000C-2950
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Sequence 16151, Application US/09949016

Patent No. 6812339

GENERAL INPORMATION:

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-20

PRIOR PILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

NUMBER OF SEQ ID NOS: 207012

SOCTWARE: PRESENCE for Windows Version 4.0

SEQ ID NO 16151

LENGTH: 146401
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APPLICANT: Ficht, Thomas A.
APPLICANT: Sowa, Blair A.
APPLICANT: Adams, L. Gary
TITLE OF INVENTION: NOVEL PROBES AND METHOD FOR IDENTIFYING
TITLE OF INVENTION: SPECIES AND BIOVARS OF BRUCELLA
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: PRAVEL, GAMBRELL, HEWITT, & KRIEGER
  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER REABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/972,791
FILING DATE: 19921106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: PRAVEL, GAMBRELL, HEWITT, & KRIEGER STREET: 1177 West Loop South, 10th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              80.0%; Score 14.4; DB 4;
93.8%; Pred. No. 3.9e+02;
iive 0; Mismatches 1;
  Mismatches
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; Patent No. 5348857
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NAME: Kettelberger, Denise M.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-850-0909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89697 rraarrcccrrrccrc 89712
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Best Local Similarity 93.0.
Local 15; Conservative
15; Conservative
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COUNTRY: USA
                                                                                                                                                                                        US-09-949-016-16151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-949-016-16151
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US-07-972-791-3/c
  Matches
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Sequence 1367, Application US/09134001C

Sequence 1367, Application US/09134001C

GENERAL INPORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS

TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: GTC-007

FILE REFERENCE: GTC-007

CURRENT APPLICATION NUMBER: US 60/064,964

PRIOR APPLICATION NUMBER: US 60/064,964

PRIOR APPLICATION NUMBER: US 60/055,779

PRIOR FILING DATE: 1997-08-14

NUMBER OF SEQ ID NOS: 5674

LENGTH: 963
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Patent No. 6703492

GENERAL INFORMATION:
APPLICANT: KINMERLY, WILLIAM JOHN

TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279

CURRENT FILING DATE: 2000-11-09

RIOR FILING DATE: 1999-11-09

NUMBER OF SEQ ID NOS: 4472

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 3819

LENGTH: 2815
                                                                                                                                                                                                                                      Gaps
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                                                          FEATURE:
COTHER INFORMATION: Description of Artificial Sequence: synthetic
CHER INFORMATION: nucleic acid sequence
US-09-710-279-3059
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                                                                                                                                                                                   Length 939;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 80.0%; Score 14.4; DB 3; Length 963; Best Local Similarity 93.8%; Pred. No. 2.5e+02; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                   Query Match

80.0%; Score 14.4; DB 4;
Best Local Similarity 93.8%; Pred. No. 2.5e+02;
Matches 15; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Staphylococcus epidermidis
                                                                                                                                                                                                                                                                                                         616 TTAATTCACGTTCCTC 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         640 rraarreacerrecre 625
              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 TTAATTCGCGTTCCTC 18
                                                                                                                                                                                                                                                                                3 TTAATTCGCGTTCCTC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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1487 GTTAATTCGCGTTC 1474
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                                                                                                                                                                                                                                                                                                                                                                                      STATE: Texas
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                  77027
                                                                      RESULT 12
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Sequence 7, Application US/07972791

Sequence 7, Application US/07972791

GENERAL INFORMATION:
APPLICANT: Ficht, Thomas A.
APPLICANT: Sowa, Blair A.
APPLICANT: Adams, L. Gary
ITLLE OF INVENTION: NOVEL PROBES AND METHOD FOR IDENTIFYING
TITLE OF INVENTION: SPECIES AND BIOVARS OF BRUCELLA
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: PRAVEL, GAMBRELL, HEMITT, & KRIEGER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 77.8%; Score 14; DB 1; Length 3345; Best Local Similarity 100.0%; Pred. No. 4.6e+02; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                 Query Match 77.8%; Score 14; DB 1; Length 3208; Best Local Similarity 100.0%; Pred. No. 4.5e+02; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: DatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/972,791

FILING DATE: 19921106

CLASSIFICATION: 4355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: PRAVEL, GAMBRELL, HEWITT, & KRIEGER
1177 West Loop South, 10th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Brucella melitensis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY AGENT INFORMATION:
NAME: Kettelberger, Denise M.
TELECHMUNICATION INFORMATION:
TELEPHONE: 713-850-0909
TELEFAX: 713-850-0165
                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: CDNA
ORIGINAL SOURCE:
CRGANISM: Brucella abortus
STRAIN: biovar 1 ($2308)
US-07-972-791-3
                                                                                                                                                                                                                                                                                                                                                                                                              1488 GTTAATTCGCGTTC 1475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 713-850-0165
TELEX: 792026
INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
LENGTH: 3345 base pairs
TELEX: 792026
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 3208 base_pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 3345 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                    2 GTTAATTCGCGTTC 15
                                                                  LENGTH: 3208 base pai
TYPE: NUCLEIC ACID
STRANDEDNESS: single
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STRAIN: biovar 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 11
US-07-972-791-7/c
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2 GTTAATTCGCGTTC 15

Gaps

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GENERAL INFORMATION:
APPLICANT: Ficht, Thomas A.
APPLICANT: Sowa, Blair A.
APPLICANT: Adams, L. Garlar A.
APPLICANT: Adams, L. Garlar A.
TITLE OF INVENITION: NOVEL PROBES AND METHOD FOR IDENTIFYING
TITLE OF INVENITION: SPECIES AND BIOVARS OF BRUCELLA
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: PRADEL, GAMBRELL, HEWITT, & KRIEGER
STREET: 1177 West Loop South, 10th Floor
                                                                                                                                                                                                            Query Match 77.8%; Score 14; DB 1; Length 3347; Best Local Similarity 100.0%; Pred. No. 4.6e+02; Matches 14; Conservative 0; Mismatches 0; Indel8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CUNTRY: Texas

COUNTRY: Dexas

COUNTRY: USA

ZIP: 77027

COMPUTER TEXADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/972,791
FILING DATE: 19921106
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kettelberger, Denise M.
TELEPHONE: 711-850.0909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Search completed: October 23, 2005, 16:07:08 Job time: 11.7318 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 6, Application US/07972791 Patent No. 5348857
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; TYPE: NUCLEIC ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: CDNA
; ORGANISM: Brucells Buis
; STRAIN: biovar 1
US-07-972-791-8
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TELEX: 792026
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 3361 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: CDNA
ORIGINAL SOURCE:
GRGANISM: Brucella ovis
US-07-972-791-6
                                                                                                                                                                                                                                                                                                            2 GTTAATTCGCGTTC 15
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US-07-972-791-6/c
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Patent No. 5348857
GENERAL INFORMATION:
APPLICANT: Ficht, Thomas A.
APPLICANT: Sowa, Blair A.
APPLICANT: Adams, L. Gary
TITLE OF INVENTION: NOVEL PROBES AND METHOD FOR IDENTIFYING
TITLE OF INVENTION: SPECIES AND BIOVARS OF BRUCELLA
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSE: PRAVEL, GAMBRELL, HEWITT, & KRIEGER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 77.8%; Score 14; DB 1; Length 3347;
100.0%; Pred. No. 4.6e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSE: PRAVEL, GAMBRELL, HEWITT, & KRIEGER STREET: 1177 West Loop South, 10th Floor CITY: Houston STATE: Texas COUNTRY: USA ZIP: 77027 COMPUTER: USA ZIP: 77027 COMPUTER: ISP PLOOPY GISK COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.25 CURRENT APPLICATION DATA: US/07/972,791 FILING DATE: 19921106 CLASSIFICATION: 435
                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
AURENT APPLICATION DATA:
PILING DATE: 19921106
CLASSIFICATION: 435
                                                                                                                                                                                                               CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kettelberger, Denise M.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-850-0909
TELEFAX: 713-850-0165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kettelberger, Denise M.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-850-0909
TELEFAX: 713-850-0165
                                                                                                                                                                                                                                                                                                                             TELEFAX: 713-850-0165
TELEX: 792026
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 3347 base pairs
TYPE: NUCLEIC ACID
STRANDEDRESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Brucella abortus
STRAIN: biovar 5
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TELES: 792026
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
IRNGTH: 3347 base pairs
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Matches 14; Conservative
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US-07-972-791-8/c
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9772231 segs, 4124568258 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                    OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                   US-10-046-955-50
18
1 cgttaattcgcgttcctc 18
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seq length: 200000000
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Maximum DB
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                                                                                                                                                                                                          Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Appl	645, A	646, A	23645, A	646, A	93618, A	978, App
ជ	25	23	23	23	23	6	97
Description	Sequence	Sequence	Sequence 23646, A	Sequence	Sequence	Sequence	Sequence
Q	US-10-046-955-50	US-10-363-345A-23645	US-10-363-345A-23646	US-10-363-483A-23645	US-10-363-483A-23646	US-10-437-963-93618	US-10-282-122A-978
я	15	20	20	71	. 12	19	. 11
Query Match Length DB ID	18	538	538	538	538	4461	127
Query Match	100.0	85.6	85.6	85.6	85.6	85.6	83.3
Score	18	15.4	15.4	15.4	15.4	15.4	15
ult No.	-	0	٣	4	2	9	7
Result No.	!		U		υ	U	υ

	equence 1074, A equence 1074, A equence 1074, A equence 81762, equence 103, Ap	Sequence 16178, A Sequence 14681, A Sequence 14682, A Sequence 14681, A Sequence 14682, A	sequence	Sequence 6603, Ap Sequence 400752, Sequence 4913, Ap Sequence 4914, Ap Sequence 4914, Ap Sequence 4914, Ap Sequence 4914, Ap
-su su su su su su su su su su su su su s	US-10-363-345 US-10-363-483 US-10-363-483 US-10-437-963 US-10-255-536 US-10-425-115	US-11-097-143-16178 US-10-363-345A-14681 US-10-363-345A-14682 US-10-363-483A-14681 US-10-363-463A-14681	US-10 US-10 US-11 US-11 US-11 US-11 US-11 US-11	US-09-974-300-6603 US-09-925-065A-400752 US-09-925-065A-18369 US-10-363-345A-4913 US-10-363-483A-4914 US-10-363-483A-4914 US-10-363-483A-4914 US-10-029-386-7594
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128 250 250 1467 9212 262 438	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	1200 1231 1231 1231 1231	230 230 3135 3257 4491 5864 15428 2731748 65 230	30 0 4 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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ALIGNMENTS

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Sequence 50, Application US/10046955;
Sequence 50, Application US/10046955;
Publication No. US20030129600A1
GENERAL INFORMATION:
APPLICANT: Secretary of the United States of America, as Represented by the APPLICANT: Control and Prevention
APPLICANT: Control and Prevention
APPLICANT: Control and Prevention
APPLICANT: Reiss, Errol
APPLICANT: Reiss, Errol
APPLICANT: Applicant Morisor Christine J.
APPLICANT: Applicant Morisor Christine J.
APPLICANT: Applicant Morisor Christine J.
APPLICANT: Control on Soo
ITILE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and
TITLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and
TITLE OF INVENTION: NUMBER: US/10/046,955
CURRENT FILING DATE: 2002-06-04
PRIOR FILING DATE: 2000-06-27
PRIOR FILING DATE: 1998-05-01
PRIOR FILING DATE: 1998-05-01
PRIOR FILING DATE: 1998-05-01
PRIOR FILING DATE: 1997-05-02
NUMBER OF SEQ ID NOS: 61
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA

ORGANISM: Fusarium oxysporum
US-10-046-955-50
              US-10-046-955-50
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Sequence 93618, Application US/10437963; Publication No. US20040123343A1; GENERAL INFORMATION: APPLICANT: La Rosa, Thomas J.; APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-437-963-93618/c
             RESULT 4
US-10-363-483A-23645
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                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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Publication No. US20040239960A1
GENERAL INFORMATION:
APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
TITLE OF INVENTION: Cytosines in genomic DNA in the sequence context of 5'-CpG-3;
FILE REPRENCE: ED1/1227
CURRENT APPLICATION NUMBER: US/10/363,345A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
SEQ ID NO 23646
                                                                                                                                                                                                                                                                                 Sequence 23645, Application US/10363345A
Publication No. US20040234960A1
GENERAL INFORMATION:
APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
APPLICANT: Alexander Ole Christian Piepenbrock
APPLICANT: Alexander Ole Christian Piepenbrock
TITLE OF INVENTION: Cytosines in genomic DNA in the sequence context of 5'-CpG-3;
FILE REPRENDER CE: E01/1227
CURRENT APPLICATION NUMBER: US/10/363,345A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
SEQ ID NO 23645
LENGTH: 538
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US-10-363-345A-23645
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US-10-363-345A-23646
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             DB 15; Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                            0; Indels
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94.1%; Pred. No. 3.3e+02;
tive 0; Mismatches 1;
          100.0%; Score 18; DB 1 100.0%; Pred. No. 9.5;
                                                            0; Mismatches
                                                                                                                                        1 CGTTAATTCGCGTTCCTC 18
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Query Match
Best Local Similarity 100.
Matches 18; Conservative
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Matches 16; Conservative
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US-10-363-345A-23646/c
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Publication No. US2005064401A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Alexander Olek

APPLICANT: Kurt Berlin

TITLE OF INVENTION: Diagnosis of illnesses or predisposition to certain

TITLE OF INVENTION: 111nesses

FILE REPERENCE: 82011

CURRENT APPLICATION NUMBER: US/10/363,483A

CURRENT PILING DATE: 2003-03

NUMBER OF SEQ ID NOS: 40712

SEQ ID NO 23646

LENGTH: 538
                                                                            APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
APPLICANT: Christian Piepenbrock
APPLICANT: Kurt Berlin
TITLE OF INVENTION: Diagnosis of illnesses or predisposition to certain
TITLE OF INVENTION: illnesses
FILE REPERENCE: 82011
CURRENT APPLICATION NUMBER: US/10/363,483A
CURRENT FILING DATE: 2003-03
SEQ ID NO.23645
LENGTH: 538
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US-10-363-483A-23646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens) OTHER INFORMATION: CpG-island No: 23645
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 538;
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Best Local Similarity 94.1%; Pred. No. 3.3e+02;
Matches 16; Conservative 0; Mismatches 1;
Sequence 23645, Application US/10363483A Publication No. US20050064401A1 GENERAL INFORMATION:
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US-10-363-483A-23646/c
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Gaps

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Score 15; DB 17; Length 127;
Pred. No. 4.9e+02;
0; Mismatches 0; Indels
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APPLICANT: AYSKING, JOHN D.
APPLICANT: Trawick, John D.
APPLICANT: Trawick, John D.
APPLICANT: Tarwick, John D.
APPLICANT: Yamanco, Robert T.
APPLICANT: Yamanco, Robert T.
TITLE OF INVENTION: Identification of Essential Genes in TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION WUMBER: 60/191, 078
FRIOR FILING DATE: 2000-03-21
FRIOR FILING DATE: 2000-03-23
FRIOR FILING DATE: 2000-03-23
FRIOR FILING DATE: 2000-03-23
FRIOR APPLICATION WUMBER: 60/206, 848
FRIOR FILING DATE: 2000-11-27
FRIOR FILING DATE: 2000-11-27
FRIOR FILING DATE: 2000-11-27
FRIOR FILING DATE: 2000-11-27
FRIOR APPLICATION WUMBER: 60/253, 625
FRIOR FILING DATE: 2000-11-27
FRIOR APPLICATION NUMBER: 60/253, 331
FRIOR FILING DATE: 2000-11-27
FRIOR APPLICATION NUMBER: 60/253, 331
FRIOR FILING DATE: 2000-11-27
FRIOR APPLICATION NUMBER: 60/259, 308
FRIOR FILING DATE: 2000-11-27
FRIOR APPLICATION NUMBER: 60/269, 308
FRIOR FILING DATE: 2000-12-22
FRIOR FILING DATE: 2000-02-16
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Pred. No. 4.9e+02;
0; Mismatches 0;
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 996
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Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari L.
APPLICANT: Zyskind, Undith W.
APPLICANT: Trawick, John D.
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 996, Application US/09815242
Patent No. US20020061569A1
GENERAL INFORMATION:
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100.0%; Pre
0;
                                                                                                                               Query Match 83.3%; Sc
Best Local Similarity 100.0%; P
Matches 15; Conservative 0;
                              ; TYPE: DNA
; ORGANISM: Enterococcus faecalis
US-10-282-122A-978
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                                                                                                                                                                                                                                                                                  117 GTTAATTCGCGTTCC 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari L.
APPLICANT: Zyskind, Judith W.
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Best Local Similarity 100.
Matches 15; Conservative
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  LENGTH: 127
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)8
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
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APPLICANT: Trawick, John
APPLICANT: Vanamoto, Robert
APPLICANT: Yanamoto, Robert
APPLICANT: Yu, H.
TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
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Pred. No. 4e+02;
0; Mismatches 1;
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CTHER INFORMATION: Clone ID: PAT_MRT4530_91986C.1

US-10-437-963-93618
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CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-03-21
PRIOR PILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR PILING DATE: 2000-09-09
PRIOR PELING DATE: 2000-10-23
PRIOR PELING DATE: 2000-10-23
PRIOR PELICATION NUMBER: 60/242,578
PRIOR PELICATION NUMBER: 60/245,578
PRIOR PELICATION NUMBER: 60/245,578
PRIOR PILING DATE: 2000-11-27
PRIOR PILING DATE: 2000-11-27
PRIOR PILING DATE: 2000-12-22
PRIOR PILING DATE: 2000-12-22
PRIOR PILING DATE: 2000-12-22
PRIOR PILING DATE: 2001-02-09
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Sequence 978, Application US/10282122A
Publication No. US20040029129A1
GENERAL INFORMATION:
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Cao, Yongwei
Wu, Wei
Boukharov, Andrey A.
Barbazuk, Brad
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APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
APPLICANT: Haeelbeck, Robert
APPLICANT: Obleen, Kari
APPLICANT: Zyskind, Judith
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Best Local Similarity 94.1%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                        SEQ ID NO 93618
LENGTH: 4461
                                                 APPLICANT
                                                                             APPLICANT
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; TYPE: DNA
; ORGANISM: Enterococcus faecalis
US-10-282-122A-531
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 GTTAATTCGCGTTCC 16
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Carr, Grant
Yamamoto, Robert
Forsyth, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Wang, Liangsu
APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
APPLICANT: Haselbeck, Robert
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Zyskind, Judith
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Trawick, Joh
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Best Local Similarity
Matches 15; Conserv
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US-10-282-122A-6332
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APPLICANT:
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TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REPERENCE: ELITRA.034

CURRENT APPLICATION NUMBER: US/10/282,122A

CURRENT PILING DATE: 2003-02-20

PRIOR PAPLICATION NUMBER: 60/201,078

PRIOR PILING DATE: 2000-03-21

PRIOR FILING DATE: 2000-05-23

PRIOR PILING DATE: 2000-05-23

PRIOR PILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-05-66

PRIOR PILING DATE: 2000-09-06

PRIOR PILING DATE: 2000-09-09

PRIOR PILING DATE: 2000-09-09

PRIOR PILING DATE: 2000-09-09

PRIOR PILING DATE: 2000-09-09

PRIOR PILING DATE: 2000-09-03

PRIOR PILING DATE: 2000-09-03

PRIOR PILING DATE: 2000-09-03

PRIOR PILING DATE: 2000-09-03
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0
APPLICANT: Carr, Grant J.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Yau, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Dentification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR PILING DATE: 2000-03-23
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-27
PRIOR PILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR PRILING DATE: 2000-11-27
PRIOR PRILING DATE: 2000-11-27
PRIOR PRILING DATE: 2000-11-27
PRIOR FILING DATE: 2000-11-27
PRIOR FILING DATE: 2001-02-16
PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 14110
SOFTWARE: PastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83.3%; Score 15; DB 9; Length 250; 100.0%; Pred. No. 5.2e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 531, Application US/10282122A Publication No. US20040029129A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Enterococcus faecalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 GTTAATTCGCGTTCC 16
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APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari
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Wall, Daniel
Trawick, John
Carr, Grant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yamamoto, Robert
Forsyth, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.
Matches 15; Conservative
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APPLICANT:
APPLICANT:
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APPLICANT:
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APPLICANT: Xu, H.
TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REPERENCE: ELITRA-034A
CURRENT APPLICATION NUMBER: 105/10/282,122A
CURRENT FILING DATE: 2003-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
PRIOR FILING DATE: 2000-11-27
PRIOR PLICATION NUMBER: 60/257,931
PRIOR PILIG DATE: 2000-12-22
PRIOR PILIG DATE: 2000-12-29
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2001-02-06
PRIOR FILING DATE: 2001-02-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3.1
SEQ ID NO 531
LENGTH: 250
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SOFTWARE: Patentin version 3.1
SEQ ID NO 6332
LENGTH: 1467
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PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR PELICATION NUMBER: 60/206, 848
PRIOR PELICATION NUMBER: 60/207, 727
PRIOR APPLICATION NUMBER: 60/20, 335
PRIOR PELING DATE: 2000-05-06
PRIOR PELING DATE: 2000-09-09
PRIOR FILING DATE: 2000-09-09
PRIOR FILING DATE: 2000-09-09
PRIOR PELING DATE: 2000-10-27
PRIOR APPLICATION NUMBER: 60/23, 347
PRIOR APPLICATION NUMBER: 60/23, 625
PRIOR APPLICATION NUMBER: 60/23, 625
PRIOR PELING DATE: 2000-11-27
PRIOR PELING DATE: 2000-11-27
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2001-02-09
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APPLICANT: Seilhamer, Jeffrey J.
APPLICANT: Stuart, Susan G.
APPLICANT: Stuart, Susan G.
APPLICANT: Stuart, Susan G.
APPLICANT: Stuart, Susan G.
APPLICANT: Mullahy, Sara J.
APPLICANT: Sara J.
APPLICANT: Sara J.
APPLICANT: Stuart J.
APPLICANT: Stua
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Sequence 43896

Sequence 43896

Sequence 43896

Publication No. US20040031072A1

GENERAL INFORMATION:

APPLICANT: La Rosa Thomas J

APPLICANT: Cao Yongwei

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(53223) B

CURRENT APPLICATION NUMBER: US/10/424,599

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 43896

LENGTH: 438
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Publication No. US20040234960A1
GENERAL INFORMATION:
APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
APPLICANT: Christian Piepenbrock
APPLICANT: Murt Berlin
TITLE OF INVENTION: Method for determining the degree of methylation of defined
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20030040615A1 hu01153717
US-09-535-459-1371
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 18;
Pred. No. 7e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    82.2%; Score 14.8; DB 10;
88.9%; Pred. No. 6.7e+02;
tive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CGTTAATTCGCGTTCCTC 18
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Best Local Similarity 88.9°
Matches 16; Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-363-345A-1073
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Patrick J. Dillon
Steven Barash
TITLE OF INVENTION: Enterococcus faecialis Polynucleotides and Polypeptides
NUMBER OF SEQUENCES: 982
CORRESPONDENCE ADDRESS:
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                                                             Query Match 83.3%; Score 15; DB 17; Length 1467; Best Local Similarity 100.0%; Pred. No. 6e+02; Matches 15; Conservative 0; Mismatches 0; Indels C
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COMPUTER: HP Vectra 486/33
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/070,927A
FILING DATE: 04-MAY-2000
CLASSIFICATION NUMBER: 60/046,655
FILING DATE: 1997-05-16
APPLICATION NUMBER: 60/044,031
FILING DATE: 1997-05-16
APPLICATION NUMBER: 60/066,009
FILING DATE: 1997-11-14
ATTORNEY/AGENT INFORMATION:
NAME: Kenley K. Hoover
REGISTRATION NUMBER: 40,302
REFERENCE/DOCKET NUMBER: 9369
TELECOMMUNICATION INFORMATION:
NAME: Kenley K. Hoover
REGISTRATION NUMBER: 40,302
TELECOMMUNICATION INFORMATION:
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100.0%; Pred. No. 7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.0%; Pred. No. 7e+
Matches 15; Conservative 0; Mismatches
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SEQUENCE DESCRIPTION: SEQ ID NO: 334:
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US-09-535-459-1371/c
Sequence 1371, Application US/09535459
Publication No. US20030046615A1
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 334, Application US/09070927A Patent No. US20020120116A1 GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: double
                                                                                                                                                                                                                                1027 GTTAATTCGCGTTCC 1041
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                                                                                                                                                                                              2 GTTAATTCGCGTTCC 16
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Maryland COUNTRY: USA
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US-10-282-122A-6332
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US-09-070-927A-334
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TITLE OF INVENTION: cytosines in genomic DNA in the sequence context of 5'-CpG-3;
FILE REFERENCE: E01/1227
CURRENT APPLICATION NUMBER: US/10/363,345A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
SEQ ID NO 1073
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: FEATURE:
FEATURE:
CTHER INFORMATION: Chemically treated genomic DNA (Homo sapiens)
CTHER INFORMATION: CpG-island No: 1073
US-10-363-345A-1073
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Search completed: October 23, 2005, 23:48:19 Job time : 79.8251 secs

Sequence 1, Appli Sequence 1, Appli Sequence 2, Appli Sequence 1, Appli Sequence 38445, A Sequence 38447, A

Sequence 12387, A Sequence 12724, A Sequence 12725, A Sequence 12125, A Sequence 5818, Ap Sequence 122, App Sequence 122, App Sequence 1227, A Sequence 12177, A Sequence 12177, A Sequence 12177, A

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J.

OM nucleic

Run on:

Sequence:

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Sequence 51, Application US/09423233
Patent No. 6372430
Fatent No. 6372430
Fatent No. 6372430
FATENCEMATION:
APPLICANT: The Government of the United States of America as TITLE OF INVENTION: Nucleic Acids for Detecting Aspergillus Species and TITLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 03063-0314WP
CURRENT APPLICATION NUMBER: 2000-06-27
NUMBER OF SEQ ID NOS: 61
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/09423233
Betent No. 6372430
APPLICANT: The Government of the United States of America as TITLE OF INVENTION: Wucleic Acids for Detecting Aspergillus Species and TITLE OF INVENTION: Other Filamentous Fungi
FILE REFERENCE: 03063-0341MP
CURRENT APPLICATION NUMBER: US/09/423,233
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 61
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 21; DB 3; Length 319; 100.0%; Pred. No. 0.22; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 21; DB 3; Length 21; 100.0%; Pred. No. 0.15; cive 0; Mismatches 0; Indels
US-09-949-016-12387
US-09-949-016-12724
US-09-270-767-2818
US-09-270-767-18100
US-09-252-991A-6522
US-08-311-731A-122
US-09-949-016-16286
US-09-949-016-16286
US-09-949-016-16286
US-09-949-016-16286
US-09-949-016-16286
US-09-949-016-12284
US-09-138-185A-1
US-09-138-185A-1
US-09-138-185A-1
US-09-138-185A-1
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US-09-423-233-6
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Best Local Similarity
Matches 21; Conserv
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Matches 21; Conserv
                US-09-423-233-51
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US-09-423-233-6
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  LENGTH: 21
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Sequence 6, Appli
Sequence 24, Appli
Sequence 87134, A
Sequence 8, Appli
Sequence 17012, A
Sequence 17012, A
Sequence 11199, A
Sequence 11, Appli
Sequence 266, Appli
Sequence 266, Appli
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206, App
120, App
24, Appl
135, App
145735,
                                                                                         October 23, 2005, 06:56:31; Search time 12.5204 Seconds (without alignments) 2744.464 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                      GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S-09-949-016-145735
S-09-949-016-146003
S-09-949-016-146271
S-09-949-016-150850
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US-09-423-233-6
US-09-949-016-87133
US-09-949-016-87134
US-09-544-3988-8
US-09-544-3988-8
US-09-949-016-17012
US-09-949-016-17012
US-09-949-016-17012
US-08-956-171E-206
US-08-956-171E-206
US-08-311-731A-120
US-08-311-731A-135
US-08-311-731A-135
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US-09-949-016-15853
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                                                                                                                                                                                                                                           1202784 segs, 818138359 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                 nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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seq length: 200000000
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Match Length
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29555
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APPLICANT: Beck, James J.
TITLE OF INVENTION: DETECTION OF WHEAT FUNGAL PATHOGENS
TITLE OF INVENTION: USING THE POLYMERASE CHAIN REACTION
NUMBER OF SEQUENCES: 24
CORRESPONDENCES: 24
CORRESPONDENCES: No. 5827695artis Corporation Patent Department
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN: Fusarium avenaceum
INDIVIDUAL ISOLATE: 64452 and R-4045 (consensus sequence)
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: 1..30
OTHER INFORMATION: /note= "3' end of small subunit
OTHER INFORMATION: rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Research Triangle Park
STATE: NC
COUNTRY: USA
ZIP: 20779-2257
COMPUTER READABLE FORM:
MEDTUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/905,314A
FILING DATE:
CLASSIFICATION: 435
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LOCATION: 182..338
OTHER INFORMATION: /note= "5.8S rRNA gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
NAME/KEY: misc_feature
LOCATION: 31..181
OTHER INFORMATION: /note= "ITS 1"
                                                                                                                                          US-08-905-314A-24
Sequence 24, Application US/08905314A
Patent No. 5827695
GENERAL INFORMATION:
APPLICANT: Beck, James J.
                                                222 CTAACACCTCGCAACTGGAGA 242
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LOCATION: 339.7504
OTHER INFORMATION: /note= "ITS
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LOCATION: 505..561
COTHEN INFORMATION: /note= "5';
CTHER INFORMATION: rRNA gene"
US-08-905-114A-24
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ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timochy
REGISTRATION UNUMBER: 38,241
REFERENCE/DOCKET NUMBER: GGC J
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-869
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 561 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
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90.5%; Score 19; DB 1; Length 561;

Query Match

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US-09-949-016-87134/c
; Sequence 87134, Application US/09949016
; Patent No. 6812339
; GENERAL INCOFFARATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; TITLE OF INVENTION: WINDER: US/09/949,016
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR PILING DATE: 2000-10-20
; PRIOR FILING DATE: 2000-10-03
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FREESE for Windows Version 4.0
; SEQ ID NO 87134
; LEMOTH: 601
                                                                                                                                                                                                               US-09-949-016-87133/c

Sequence 87133, Application US/09949016

Sequence 87133, Application US/09949016

Sequence 87133, Application US/09949016

Sequence 87133, Application

GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOCTHARE: FastEEQ for Windows Version 4.0

SEQ ID NO 87133

LENGTH: 601
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Best Local Similarity 100.0%; Pred. No. 2.6;
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Best Local Similarity 85.7%; Pred. No. 75;
Matches 18; Conservative 0; Mismatches
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                         0; Mismatches
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                                                                      1 CTAACACCTCGCAACTGGA 19
                         Conservative
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US-09-949-016-87133
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US-09-949-016-87134
                         19;
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US-09-949-016-17012/c
; Sequence 17012, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;
; FILE REPERBENCE: CL001307
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/231,768
; PRIOR PAPLICATION NUMBER: 60/231,768
; PRIOR PELING DATE: 2000-10-03
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 2000-10-03
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FREESEQ for Windows Version 4.0
; SEQ ID NO 17012
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| Batent No. 6812339
| GENERAL INPORMATION:
| APPLICANT: VENTER, J. Craig et al. |
| APPLICANT: VENTER, J. Craig et al. |
| TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
| FILE REFERENCE: CLOOL307
| CURRENT APPLICATION NUMBER: US/09/949,016
| CURRENT PILING DATE: 2000-04-14
| PRIOR APPLICATION NUMBER: 60/241,755
| PRIOR PILING DATE: 2000-10-02
| PRIOR FILING DATE: 2000-10-03
| PRIOR FILING DATE: 2000-10-03
| PRIOR FILING DATE: 2000-0-0-09
| PRIOR FILING DATE: 2000-0-09-08
| NUMBER OF SEQ ID NOS: 207012
| SOFFWARE: FASELSEQ for Windows Version 4.0
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                                    Length 33769;
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                                                                                   3; Indels
                              Ouery Match 77.1%; Score 16.2; DB 4; Best Local Similarity 85.7%; Pred. No. 1.3e+02; Matches 18; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 77.1%; Score 16.2; DB 4; Best Local Similarity 85.7%; Pred. No. 1.5e+02; Matches 18; Conservative 0; Mismatches 3;
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; LOCATION: (1)...(89625)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-17012
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NAME/KEY: misc_feature
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ORGANISM: Human
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LENGTH: 140844
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; LOCATION: (33739),(33749),(33758)
; OTHER INFORMATION: Identity of nucleotide sequences at the above locations are unknous-09-543-771B-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: unsure

: LOCATION: (33739), (33749), (33758)

: COCATION: Identity of nucleotide sequences at the above locations are unkno

US-09-544-398B-8
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85.7%; Pred. No. 1.3e+02;
tive 0; Mismatches 3; Indels 0;
                                                                                                                                                                                            GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Carulli, John P.
APPLICANT: Carulli, John P.
APPLICANT: Little, Randall D.
APPLICANT: Little, Randall D.
APPLICANT: APPLICANT:
TITLE OF INVENTION: High bone mass gene of 11q13.3
FILE REFERENCE: 032796-013
CURRENT APPLICATION NUMBER: US/09/544,398B
CURRENT PRING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: US 60/011,449
PRIOR FILING DATE: 1999-01-13
PRIOR PLILING DATE: 1998-01-13
PRIOR PLILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 641
SOCTHWARE: FASTESEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 33769
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/09543771B
Parent No. 6780609
GENERAL INFORMATION:
APPLICANT: Carulli, John P.
APPLICANT: Little, Randall D.
APPLICANT: Recker, Robert R.
APPLICANT: Johnson, Mark L.
TITLE OF INVENTION: High bone mass gene of 11q13.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/543,771B
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/229,319
PRIOR FILING DATE: 1999-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR FILING DATE: 1998-01-3
PRIOR FILING DATE: 1998-01-3
NUMBER OF SEQ ID NOS: 641
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 33769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25831 CTAGCACCTCTCAACTGGACA 25851
66 CTAGCACCTCTCAACTGGACA 46
                                                                                                                               Sequence 8, Application US/09544398B Patent No. 6770461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 CTAACACCTCGCAACTGGAGA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 85.79
Matches 18; Conservative
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                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
US-09-543-771B-8
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; TOPOLOGY: US-08-781-986A-206
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US-09-815-923-11/c
Sequence 11, Application US/09815923
Sequence 11, Application US/09815923
GENERAL IN. 6787642
GENERAL INFORMATION:
APPLICANT: Gill, Sarjeet S.
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Use of Insect Cell Membrane Transporters as No. 6787642e1
TITLE OF INVENTION: Target Sites for Insecticides
FILE REFERENCE: 023070-093800US
GURRENT APPLICATION UNBER: US/09/815,923
CURRENT APPLICATION UNBER: 2001-03-23
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Charles Kunsch
Gil H. Choi
Patrick S. Dillon
Craig A. Rosen
Craig A. Rosen
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
CORRESPONDENCE ADDRESS:
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                                                                                           77.1%; Score 16.2; DB 4; Length 140844; 85.7%; Pred. No. 1.6e+02; tive 0; Mismatches 3; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 73.3%; Score 15.4; DB 4; Length 2034; Best Local Similarity 94.1%; Pred. No. 2.3e+02; Matches 16; Conservative 0; Mismatches 1; Indels 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage

COMPUTER: HP Vectra 486/33

OPERATING SYSTEM: MSDOS version 6.2

SOFTWARE: ASCII Text

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Human Genome Sciences, Inc
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/956,171E FILING DATE: 20-Oct-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: orphan transporter US-09-815-923-11
                                                                                                                                                                                                                                    42302 CTAGCACCTCTCAACTGGACA 42282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 11
US-08-956-171E-206/c
; Sequence 206, Application US/08956171E
; Patent No. 6593114
; GENERAL INFORMATION:
                                                                                                                                                                                        1 CTAACACCTCGCAACTGGAGA 21
; LOCATION: (1)...(140844)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-14199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           423 CACCACGCAACTGGAGA 407
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                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Manduca sexta
                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 11
LENGTH: 2034
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                         RESULT 10
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US-08-781-986A-206/C

Sequence 206, Application US/08781986A

Sequence 206, Application US/08781986A

Patent No. 6737248

GENERAL INFORMATION:
APPLICANT: Charles Kunsch
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences NUMBER OF SEQUENCES: 5255
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73.3%; Score 15.4; DB 4; Length 29555; 94.1%; Pred. No. 3.4e+02; ive 0; Mismatches 1; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
                                                                                                                                                                 NAME: Mark J. Hyman
REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB248P1
                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
;
SEQUENCE DESCRIPTION: SEQ ID NO: 206:
US-08-956-171E-206
                                                                 FILING DATE: January 5, 1996
APPLICATION WUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/009,861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/781,986A
FILING DATE:
                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (340) 314-1224
TELEFAX: (301) 309-8439
INFORMATION FOR SEQ ID NO: 206:
SEQUENCE CHARACTERISTICS:
LENGTH: 29555 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1809 TAATACCTCGCAACTGG 1793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAVE: Benson, Bob
REGISTRATION NUMBER: 30,446
REPERENCE/DOCKET NUMBER: PB2
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 206:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 TAACACCTCGCAACTGG 18
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TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 73.39
Best Local Similarity 94.1<sup>3</sup>
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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RESULT 15

US-08-311-731A-135/c

Sequence 135, Application US/08311731A

Sequence 135, Application US/08311731A

Patent No. 6583266

GENERAL INFORMATION:
APPLICANT: SMITH, DOUGLAS

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES

TITLE OF INVENTION: RELATING TO MYCOBACTERIUM TUBERCULOSIS AND LAPRAE FOR

TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS

NUMBER OF SEQUENCES: 411

CORRESPONDENCE ADDRESS:
                     RELATING TO MYCOBACTERIUM TUBERCULOSIS AND LAPRAE FOR DIAGNOSTICS AND THERAPEUTICS 411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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ZIP: 02210
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: DATA
COMPUTER: DATA
COMPATIBLE
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/311,731A
                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/311,731A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 73.3%; Score 15.4; DB 4; Best Local Similarity 94.1%; Pred. No. 3.5e+02; Matches 16; Conservative 0; Mismatches 1;
                                                                                                              E: WOLF, GREENFIELD & SACKS, P.C. 600 ATLANTIC AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WOLF, GREENFIELD & SACKS, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C0044/7125
                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION ....
FILING DATE:
CLASSIFICATION: 530
ATTORNEY AGENT INFORMATION:
NAME: GATES, EDWARD R.
REGISTRATION NUMBER: 31,616
REFERENCE/DOCKET NUMBER: C0044
TELECOMMUNICATION INFORMATION:
TELEFAX: 617/720-241
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: MYCOBACTERIUM LEPRAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23279 CACCTCGCAACTGGCGA 23263
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: circular MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 38494 base pairs
TYPE: nucleic acid
                                                                                                                                 STREET: 600 ATLANTIC
CITY: BOSTON
STATE: MASSACHUSETTS
                     TITLE OF INVENTION: REI
TITLE OF INVENTION: DIN
NUMBER OF SEQUENCES: 4:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: BOSTON
STATE: MASSACHUSETTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                           02210
                                                                                                                 ADDRESSEE:
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                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
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                                                                                                                                                                                                                                                            Sequence 120, Application US/08311731A

Batent No. 6583266

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: NUCLAIC ACID AND AMINO ACID SEQUENCES
TITLE OF INVENTION: RELATING TO MYCOBACTERIUM TUBERCULOSIS AND LAPRAE FOR TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
CORRESPONDENCES: 411

CORRESPENDENCE ADDRESSE:
ADDRESSEE: WOLF, GREENFIELD & SACKS, P.C.
STREET: 600 ATLANTIC AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                      Gaps
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vuery Match
73.3%; Score 15.4; DB 4; Length 29555;
Best Local Similarity 94.1%; Pred. No. 3.4e+02;
Matches 16; Conservative 0; Mismatchee 1. T-1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24, Application US/08311731A
Patent No. 6583266
GENERAL INFORMATION:
APPLICANT: SMITH, DOUGLAS
APPLICANT: MAO, JEN-I
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/311,731A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICALL.

APPLICALL.

FILING DATE:
CLASSIFICATION: 530
ATTORNEY AGENT INFORMATION:
NAME: GATES, EDWARD R.
REGISTRATION NUMBER: 31,616
REFERENCE/DOCKET NUMBER: C0044/7125
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/720-3500
TELEPHONE: 617/720-2441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: MYCOBACTERIUM LEPRAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELERAX: 617/720-2441
INFORMATION FOR SEQ ID NO: 120:
SEQUENCE CHARACTERISTICS:
LENGTH: 36181 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: circular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28405 AACACCGCCAACTGGA 28389
                                                                                                                                                         1809 TAATACCTCGCAACTGG 1793
                                                                                                              2 TAACACCTCGCAACTGG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 94.1
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MASSACHUSETTS: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: WOL
STREET: 600 AT
CITY: BOSTON
STATE: MASSACH
                                                                                                                                                                                                                                                   US-08-311-731A-120/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 14
US-08-311-731A-24/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-311-731A-120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                              8
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Query Match
Best Local Similarity 94.1%; Pred. No. 3.5e+02;
Matches 16; Conservative 0; Mismatches 1; Indels 0;
FILING DATE:

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

NAME: GATES, EDWARD R.

REGISTRATION NUMBER: 31,616

REFERENC/DOCKET NUMBER: 31,616

REFERENC/DOCKET NUMBER: 07044/7125

TELEPHONE: 617/720-3500

TELEPHONE: 617/720-3441

INFORMATION FOR SEQ ID NO: 135:

SEQUENCE CHARACTERISTICS:

LEMOTH: 38675 base pairs

TYPE: nucleic acid
STRANDEDNESS: double

OPOLOGY: circular

MOLECULE TYPE: DNA (genomic)

HYPOTHETICAL: NO

ANTI-SENSE: NO

ORGANISM: Mycobacterium leprae

US-08-311-731A-135

T3.3%; Score 1:
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0;

Gaps

; 0

Search completed: October 23, 2005, 16:07:15 Job time : 19.5204 secs

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October 23, 2005, 15:20:46; Search time 90.7959 Seconds (without alignments) 1907.926 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                        OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 2000000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Sequence 51, Appl	Sequence 6, Appli	Sequence 5, Appli	Sequence 5, Appli	Sequence 367816,	Sequence 367817,	Seguence 784700,
SUMMARIES	ΠD	US-10-046-955-51	US-10-046-955-6	US-10-829-661-5	US-10-672-300-5	US-09-925-065A-367816	US-09-925-065A-367817	US-11-036-317-784700
		15	15	21	21	12	12	24
	Query Match Length DB	21	319	552	552	636	636	25
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10-424- 10-027	US-10-3/4-9/9-8 US-10-182-936A-8 US-10-731-739-8 US-10-477-228A-8 US-10-477-173-8 US-10-834-377-8 US-10-834-377-8 US-10-322-281-10 US-10-323-345A-31749 US-10-363-445A-31750 US-10-363-445A-31750	-10-363-483A -10-425-115-9 -10-369-493-7 -10-425-115-1 -10-282-122A -11-097-143-1
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ALIGNMENTS

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US-10-046-955-51

US-10-046-955-51

Sequence 51, Application US/10046955

Publication No. US2003129600A1

GENERAL INFORMATION:

APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: Secretary of the Department of Health and Human Services, Centers for Di APPLICANT: Control and Prevention

APPLICANT: Reiss, Errol

APPLICANT: Reiss, Errol

APPLICANT: Reiss, Errol

APPLICANT: Reiss, Errol

APPLICANT: Roles, Christine J.

APPLICANT: Roles, Christine J.

APPLICANT: Aldorevoich, Liliana

APPLICANT: PREENCE: 6395-6206

CURRENT APPLICATION NUMBER: US 60/045,400

PRIOR FILING DATE: 1998-05-01

PRIOR FILING DATE: 1997-05-02

PRIOR FILING DATE: 1997-05-02

NUMBER OF SEQ ID NOS: 61

LENGTH: 21

LENGTH: 21

LENGTH: 21

TYPE: DAA

ORGANISM: Fusarium solani

US-10-046-955-51
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Gaps
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TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
FRIOR PAPLICATION NUMBER: US 60/243,096
FRIOR PILING DATE: 2000-11-20
FRIOR PELING DATE: 2000-11-20
FRIOR PELICATION NUMBER: US 60/250,092
FRIOR FILING DATE: 2000-11-6
FRIOR APPLICATION NUMBER: US 60/261,766
FRIOR PILING DATE: 2001-01-6
FRIOR PILING DATE: 2001-01-6
FRIOR PILING DATE: 2001-01-6
FRIOR APPLICATION NUMBER: US 60/261,766
FRIOR PILING DATE: 2001-01-6
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; Pred. No. 12; 
0; Mismatches 0; Indels
                                                                          DB 21; Length 552;
12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/10672300
; Sequence 5, Application US/10672300
; Publication No. US20050048509A1
; GENERAL INFORMATION:
; APPLICANT: HAN, XIANG-YANG
; TITLE OF INVANITION: DIAGNOSIS OF INVASIVE MOLD INFECTION
; FILE REFERENCE: UTXC: 766.72,300
; CURRENT APPLICATION NUMBER: US/10/672,300
; CURRENT FILING DATE: 2003-09-26
; PRIOR APPLICATION NUMBER: 60/414,008
; PRIOR FILING DATE: 2002-09-27
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: Patentin Ver: 2.1
; SEQ ID NO 5
LENGTH: 552
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Pred. No. 1.6e+02;
0; Mismatches 2;
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                                                                      11arity 100.0%; Score 19; 100.0%; Pred. No. Conservative 0; Mismatch
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Best Local Similarity 100.0%;

Matches 19; Conservative 0
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Best Local Similarity 90.0
Matches 18; Conservative
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US-09-925-065A-367816
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                                                                          Query Match
Best Local Similarity
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US-10-672-300-5
US-10-829-661-5
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Sequence 6, Application US/10046955

Publication No. US20030129600A1

GENERAL INFORMATION

GENERAL INFORMATION

APPLICANT: The Government of the United States of America, as Represented by the APPLICANT: Control and Prevention

APPLICANT: Control and Prevention

APPLICANT: Addorevich, Liliana

APPLICANT: Control

TITLE OF INVENTION: Other Filamentous Fungi

FILE REFERENCE: 6395-62064

CURRENT FILING DATE: 2002-06-04

FRICE REFERENCE: 6395-62064

CURRENT FILING DATE: 2000-06-27

PRIOR FILING DATE: 1998-05-01

PRIOR PLICATION NUMBER: US 60/045,400

FRIOR FILING DATE: 1999-05-02

FRIOR FILING DATE: 1997-05-02

WUMBER OF SEQ ID NOS: 61

SSOFTWARE: Patentin Ver. 2.0

SEQ ID NO 6

SEQ ID NO 6

SEQ ID NO 6

SEQ ID NOS: 61

SEQ ID NO 6

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Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 21; Conservative 0; Mismatches 0; Indels (
                        ch 100.0%; Score 21; DB 15; Length 21; 1 Similarity 100.0%; Pred. No. 0.99; 21; Conservative 0; Mismatches 0; Indels
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APPLICANT: TARRAND, JEFFREY J.
APPLICANT: PHAM, AUDREY S.
APPLICANT: PHAM, AUDREY S.
TITLE OF INVENTION DIAGNOSIS OF MOULD INFECTION
FILLS REPERENCE: UTSCHESUS
CURRENT APPLICATION NUMBER: US/10/829,661
CURRENT FILING DATE: 2004-04-21
PRIOR FILING DATE: 2004-09-26
PRIOR APPLICATION NUMBER: 10/672,300
PRIOR PLING DATE: 2002-09-26
PRIOR PLING DATE: 2002-09-26
PRIOR PLING DATE: 2002-09-27
NUMBER OF SEQ ID NOS: 29
SOFTWARE PARCELLIN Ver. 2.1
SEQ ID NO S: 29
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Publication No. US20050009051A1
GENERAL INFORMATION:
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US-10-046-955-6
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ORGANISM: Fusarium sp
                             Query Match
Best Local Similarity
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Sequence 123319, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT APPLICATION NUMBER: US/10/424,599
SURMER OF SEQ ID NOS: 285684
SEQ ID NO 123319
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US-10-027-632-136133

Sequence 136133, Application US/10027632

Publication No. US202020198371A1

GENDRAL INFORMATION:

APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide

TITLE OF INVENTION: Polymorphisms in the Human Genome

FILE REFERENCE: 108827.129

CURRENT PAPLICATION NUMBER: US 60/218,006

PRIOR APPLICATION NUMBER: US 60/218,006

PRIOR APPLICATION NUMBER: US 60/198,676

PRIOR APPLICATION NUMBER: US 60/198,676

PRIOR PELING DATE: 2000-04-20

PRIOR PELING DATE: 2000-03-24

PRIOR APPLICATION NUMBER: US 60/195,218

PRIOR APPLICATION NUMBER: US 60/195,33

PRIOR APPLICATION NUMBER: US 60/195,33

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR APPLICATION NUMBER: US 60/157,363

PRIOR PLING DATE: 1999-10-23

PRIOR FILING DATE: 1999-10-23

PRIOR FILING DATE: 1999-09-28

PRIOR FILING DATE: 1999-09-28
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Pred. No. 3.3e+02;
0; Mismatches 3;
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Pred. No. 3.2e+02;
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85.7%;
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Best Local Similarity 85.7
Matches 18; Conservative
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Best Local Similarity
Matches 18; Conserv
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TITLE OF INVENTION: Method of Analysis of Alternative Splicing in Mouse
FILE REPERBENE: 3654.1
CURRENT APPLICATION WINBER: US/11/036,317
CURRENT FILING DATE: 2005-01-13
PRIOR APPLICATION NUMBER: US 60/536,639
PRIOR FILING DATE: 2004-01-13
NUMBER OF SEQ ID NOS: 991174
SEQ ID NOS: 991174
SEQ ID NO 784700
LENGTH: 25
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APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Indentification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/25,065A
CURRENT FILING DATE: 2000-108-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR FILING DATE: 2000-11-20
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
SPRIOR FILING DATE: 200
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Publication No. US20050228172A9
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US-11-036-317-784700/c
; Sequence 784700, Application US/11036317
; Publication No. US20050214823A1
; GENERAL INPORMATION:
; APPLICANT: Williams, Alan
   230 CTAAAACCTCTCAACTGGAG 211
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Matches 18; Conservative
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; ORGANISM: Mus musculus
US-11-036-317-784700
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Best Local Similarity
Matches 18; Conserv
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, ORGANISM: Human
US-10-027-632-136133
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US-10-027-632-136134
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APPLICANT: WANG, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
ITTLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US 60/218,006

PRIOR APPLICATION NUMBER: US 60/218,006

PRIOR FILING DATE: 2000-07-12

PRIOR PLING DATE: 2000-07-22

PRIOR PLING DATE: 2000-03-29

PRIOR PLING DATE: 2000-03-29

PRIOR PLING DATE: 2000-03-29

PRIOR PLING DATE: 1999-11-23

PRIOR PLING DATE: 1999-11-23

PRIOR FILING DATE: 1999-09-28

PRIOR FILING DATE: 1999-08-09

PRIOR PRIOR DATE: 1999-08-09
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JAPPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide

TITLE OF INVENTION: Polymorphisms in the Human Genome

FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US/10/27,632

CURRENT FILING DATE: 2002-04-30

PRIOR APPLICATION NUMBER: US 60/198,676

PRIOR APPLICATION NUMBER: US 60/198,676

PRIOR APPLICATION NUMBER: US 60/193,483

PRIOR APPLICATION NUMBER: US 60/193,483

PRIOR FILING DATE: 2000-04-20

PRIOR FILING DATE: 2000-03-29

PRIOR FILING DATE: 2000-02-24

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-09-28

PRIOR APPLICATION NUMBER: US 60/166,358

PRIOR FILING DATE: 1999-09-28

PRIOR APPLICATION NUMBER: US 60/146,002

PRIOR APPLICATION NUMBER: US 60/146,002
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SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 136133
LENGTH: 499
                                                                                               ; Sequence 136134, Application US/10027632; Publication No. US20020198371A1
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; Publication No. US20030204075A9
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US-10-027-632-136133
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US-10-027-632-136134
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APPLICANT' Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 10827.129

CURRENT PILIAGENERICE: 2002-04-30

PRIOR PAPLICATION NUMBER: US 60/218,006

PRIOR PLICATION NUMBER: US 60/198,676

PRIOR FILING DATE: 2000-07-12

PRIOR FILING DATE: 2000-04-20

PRIOR FILING DATE: 2000-04-20

PRIOR FILING DATE: 2000-03-24

PRIOR FILING DATE: 2000-03-24

PRIOR FILING DATE: 1999-11-23

PRIOR PLICATION NUMBER: US 60/165,363

PRIOR PLICATION NUMBER: US 60/166,363

PRIOR PLICATION NUMBER: US 60/166,363

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-109-08

PRIOR FILING DATE: 1999-09-08

PRIOR FILING DATE: 1999-09-08

PRIOR FILING DATE: 1999-08-09

PRIOR FILING DATE: 1999-08-09

SOFTWARE: FRASERGE for Windows Version 4.0

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LENGTH: AVS
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TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Indentification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR FILING DATE: 2000-10-24
PRIOR FILING DATE: 2000-11-20
PRIOR FILING DATE: 2000-11-20
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2000-11-30
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85.7%; Pred. No. 3.3e+02;
iive 0; Mismatches 3;
Query Match 77.1%; Score 16.2; DB 17; Best Local Similarity 85.7%; Pred. No. 3.3e+02; Matches 18; Conservative 0; Mismatches 3;
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Publication No. US20030204075A9
GENERAL INFORMATION:
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; Publication No. US20050228172A9
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT PILING DATE: 2001-08-08
FRIOR APPLICATION NUMBER: US 60/243,096
FRIOR FILING DATE: 2000-10-24
FRIOR FILING DATE: 2000-11-20
FRIOR FILING DATE: 2000-11-30
FRIOR FILING DATE: 2001-11-30
FRIOR FILING DATE: 2001-01-46
FRIOR FILING DATE: 2001-01-66
FRIOR APPLICATION NUMBER: US 60/261,766
FRIOR APPLICATION NUMBER: US 60/280,846
FRIOR APPLICATION NUMBER: US 60/2
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85.7%; Pred. No. 3.3e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                              Query Match 77.1%; Score 16.2; DB 12; Best Local Similarity 85.7%; Pred. No. 3.3e+02; Matches 18; Conservative 0; Mismatches 3;
PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: US 60/289,846
PRIOR FILING DATE: 2001-05-09
NUMBER OF SEQ ID NOS: 957086
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 487968
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Publication No. US20040123343A1
GENERAL INFORMATION:
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 85.7
Matches 18; Conservative
                                                                                                                                                                                                                                                  TYPE: DNA
CORGANISM: Homo sapiens
US-09-925-065A-487968
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-925-065A-188598/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-925-065A-188598
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APPLICANT:
APPLICANT:
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| FILE REFERENCE: 38-21(53221)B
| CURRENT APPLICATION WUMBER: US/10/437,963
| CURRENT FILING DATE: 2003-05-14
| NUMBER OF SEQ ID NOS: 204966
| SEQ ID NO 41403
| TYPE: DNA
| ORGANISM: Oryza sativa
| ORGANISM: Oryza sativa
| ORGANISM: Oryza sativa
| OTHER INFORMATION: Clone ID: PAT_MRT4530_44751C.1
| OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER
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